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Result
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          otal number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    on:
    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                295
112.5
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US-09-766-396-26
554
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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                                        665
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RIMSSI
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A28968
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JC6166
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S67489
JC6167
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                             ferrichrome transp
somatostatin II pr
WD repeat domain p
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hypothetical prote
transcription acti
membrane protein B
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cortistatin precur
somatostatin precu
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somatostatin II pre
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somatostatin, panc
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Cortistatin precursor - rat
C;Species: Rattus norvegicus (Norway rat)
C;Becies: Rattus norvegicus (Norway rat)
C;Date: 28-Oct-1996 #sequence_revision 27-Feb-1997 #text_change 16-Jul-1999
C;Accession: S67489
C;Accession: S67489
R;de Lecea, L; Criado, J.R.; Prospero-Garcia, O.; Gautvik, K.M.; Schweitzer, P.; Danlarture 381, 242-245, 1996
A;Title: A cortical neuropeptide with neuronal depressant and sleep-modulating proper A;Reference number: S67489; MUID:96208649; PMID:8622767
A;Accession: S67489
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-112 CDEL>
A;Cross-references: EMBL:U51919; NID:g1335909; PIDN:AAC52585.1; PID:g1335910
C;Superfamily: somatostatin

Query Match Best Local Similarity

53.2%;

Score 295; DB 2; Pred. No. 1.1e-23;

Length 112;

RESULT 2 \$67489 cortistatin

B 8

Qy	Query Match Best Local Similarity Matches 105; Conser	C;Comment: This protein binds to somate C;Superfamily: somatostatin F;1-20/Domain: signal sequence #status F;97-100/Region: somatostatin receptor	A; Molecule type: mRNA A; Residues: 1-105 <fuk> A; Cross-references: DDBJ: AB000263; A; Experimental source: brain</fuk>	R; Fukusumi, S.; Kit. Biochem. Biophys. R A; Title: Identifica A; Reference number: A; Accession: JC5414	UC34.4 cortistatin-like protein C; Species: Homo sapiens C; Date: 10-Jun-1997 #sec C; Accession: JC5414	RESULT 1			ωN					ω ω 4-ω		30 31	
1 MP	ch 1 Sim 105;	This lly: s iin: s egion:	type: 1-10 Eerenc	S.; lophys lentif enumb numb	n-like Homo Jun-1	à.		67 66.5	67.5 67	6 8 8	68.5	69	60	69.5 69	70	70	
LSPGLLLI SPGLLLI	100 ilarity 100 Conservative	protein comatosta signal se somatos	mRNA)5 <fuk>)es: DDBJ ;ource: h</fuk>	mi, S.; Kitada, Biophys. Res. (Identification nce number: JC54 ion: JC5414	protein precursor sapiens (man) .997 #sequence_revi			2221	12.2					12.5 12.5		12.6	
LLSGAI	100.0%; 100.0%; ative	binds tin equence	r:AB000	C.; Ta Commun and cl	n precu (man) quence			462 233	374	950	/4 1493	664	535 535	882 252	1801	14	
PAT	.09	re# to	026	har JID	_re			221	N N	N N	2 4	N I	- 2	N N	N	งผ	
ALPLEGGPTGI ALPLEGGPTGI	,0	to somatostat: #status pred: receptor bind:	; NID:g205	, C.; Takekawa, S.; Kizawa, Commun. 232, 157-163, 1997 n and characterization of a 5414; MUID:97236300; PMID:9	otein precursor – human iens (man) #sequence_revision 18-Jul-1997		ALIGNMENTS	JC2462 T36942	F87280	H82182 A82986	T10757	D83231	G97497	H82754 AD2716	T26774	A60840	
MPLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA 	DB 2; 1e-51; 0;	to somatostatin receptors #status predicted <sig> receptor binding #status</sig>	NID:g2055231; PIDN:BAA19770.1; PID:g2055232	H.; nove 12512	ıl-1997 #te		NTS										
IRKSSLLTFI	Length 105; Indels	on a predic	BAA19770.1		<pre>#text_change</pre>			gastric : probable	este	conse	MAP I	hypo	ribo	fami. ribo	hypot	somat	
AWWFEWTSQ	0; Gaps	pituitary GH4 ted	l; PID:g20	<pre>akamoto, J.; Miyamoto, ! human cortistatin-like .</pre>	20-Jun-2000			gastric inhibitory probable DNA-bindi	esterase A [import	conserved hypothet adenvlate cyclase	MAP kinase kinase	hypothetical prote	ribonuclease p	family 3 glycoside ribonuclease [impo	hypothetical p		
A 60 A 60	0;	4 cell and	55232	oto, M.; Hin -like peptid	00			tory	port	thet	ase	rote	precu	[impo	prote	, , n H	
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C;Comment: This protein a C;Genetics:
A;Gene: pss2
C;Superfamily: somatostat
C;Keywords: brain; hormon
                                             C;Reywords: hormone; neuropeptide
E;1-24/Domain: signal sequence #status predicted <SIG>E;25-88/Domain: propeptide #status predicted <PRO>E;25-88/Domain: propeptide #status predicted <PRO>E;25-88/Product: somatostatin-28 #status predicted <M28>E;103-116/Product: somatostatin-14 #status
                                                                                                                                                           A;Cross-references: GB:M31217; NID:g163636; PIDN:AAA30744.1; PID:g163637 A;NOte: the authors translated the codon ATT for residue 65 as Asn C;Comment: Somatostatin inhibits the release of somatotropin. C;Superfamily: somatostatin
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A;Residues: 1-103 <TOS>
A;Cross-references: GB:U68137; NID:g1890652; PIDN:AAC60094.1; PID:g1890653
A;Cross-references: GB:U68137; NID:g1890652; PIDN:AAC60094.1; PID:g1890653
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                                                                                                                                                                                                                                                                            A; Status: preliminary A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                          A; Reference number: A40929; A; Accession: A40929
                                                                                                                                                                                                                                                                                                                                               Mol. Endocrinol. 2, 209-216, 1988
A;TITIE: Structure and evolution of somatostatin genes.
A;Reference number: A40929; MUID:88288237; PMID:2899837
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C;Species: Rana ridibunda (laughing frog)
C;Date: 11-Apr-1997 #sequence_revision 09-May-1997 #text_change 16-Jul-1999
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                          F;105-116/Disulfide
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R; Su, C.J.; White, J.W.; Li,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Tostivint, H.; Lihrmann, I.; Bucharles, C.; Vieau, D.; Coulouarn, Y.; Fournier, A. Proc. Natl. Acad. Sci. U.S.A. 93, 12605-12610, 1996
A;Title:—Focurrence of two somatostatin variants in the frog brain: Characterization;Reference number: JC6166; MUID:97057290; PMID:8901629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Accession: JC6167
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Best Local S
Matches 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Species: Bos primigenius taurus (cattle)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contains:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Alternate names:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gene: pss2
Superfamily: somatostatin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           names: preprosomatostatin somatostatin 28 (SS-28) somatostatin 14 (SS-14); somatostatin 28 (SS-28)
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                                                                                                                                                                                                                                                                                                                                                                                                                   W.H.; Luo,
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                                                                                                                                                                                                                                                                                                                                                                                                                   C.C.; Frazier, M.L.;
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                     A; Molecule type: mRNA
A; Residues: 1-116 < RE
                                                                A; Status: translated from
                                                                                        A; Reference number: I51829;
A; Accession: I51829
                                                                                                                                                                                                                          A; Residues: 38-78, 'H', 80-116 < RES>
                                                                                                                                                                                                                                                                       A; Status: translated from GB/EMBL/DDBJ
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Science 238, 1126-1129,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    J. Biol. Chem. 258, 8788-8793, 1983
A; Title: Nucleotide and amino acid sequence
A; Reference number: A01431; MUID:83238516; F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N; Contains: somatostatin-14; somatostatin C; Species: Rattus norvegicus (Norway rat)
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A; Residues: 1-116 < ARG>
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A; Residues: 1-116 < MON>
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Best Local S
Matches 34
                       -116 <RE2>
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GB:M25890; NID:g207030;
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1987
                                                                   GB/EMBL/DDBJ
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31.5%; Pred.
                                                                                                              MUID:85303584;
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R;Tavianini, M.A.; Hayes, T.E.; Magazin, M.D.; Minth, C.D.; Dixon, J. Biol. Chem. 259, 11798-11803, 1948.
A;Title: Isolation, characterization, and DNA sequence of the rat sA;Reference number: A22529; MUID:85006903; PMID:6148343
A;Accession: A22529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: GB:K02248; NID:g207014; PIDN:AAA42161.1; R;Goodman, R.H.; Jacobs, J.W.; Dee, P.C.; Habener, J.F. J. Biol. Chem. 257, 1156-1159, 1982
A;Title: Somatostatin-28 encoded in a cloned cDNA obtained fi A;Reference number: I55220; MUID:82120034; PMID:6120163
                                                                                                                                                                                                                A;Cross-references: GB:J00788; NID:g207018; PIDI R;Goodman, R.H.; Montminy, M.R.; Low, M.J.; Habs Adv. Exp. Med. Biol. 188, 31-47, 1985
A;Title: Biosynthesis of rat preprosomatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Title: A new prosomatostatin-derived peptide reveals a pattern for prohormone cleav A;Reference number: A47598; MUID:88070564; PMID:2891188
A;Accession: A47598
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A;Note: the authors translated the codon ACC for residue 43 as Tyr
R;Argos, P.; Taylor, W.L.; Minth, C.D.; Dixon, J.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Date: 03-Aug-1984 *sequence_revision 03-Aug-1984 *text_change 18-Jun-1999 C;Accession: A20983; A01431; A47598; A22529; I55220; I51829 R;Montminy, M.R.; Goodman, R.H.; Horovitch, S.J.; Habener, J.F. Proc. Natl. Acad. Sci. U.S.A. 81, 3337-3340, 1984 A;Title: Primary structure of the gene encoding rat preprosomatostatin. A;Reference number: A20983; MUID:84221954; PMID:6145156
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   PIDN: AAA42167.1;
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Habener, J.F.
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C; Comment: Somat
C; Genetics:
A; Introns: 46/3
C; Function:
somatostatin precursor - chicken c;Species: Gallus gallus (chicken) C;Species: Gallus gallus (chicken) C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999 C;Accession: S20630 R;Nata, K.; Kobayashi, T.; Karahashi, K.; Kato, S.; Yamamoto, H.; Yonekura,
                                                                                                   RESULT
S20630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N;Contains: somatostatin 14 (SS-14); somatostatin C;Species: Mus musculus (house mouse) C;Date: 31-pec-1990 #sequence_revision 31-Jan-1997 C;Accession: S08416
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3; Keywords: hormone; neuropeptide
6; 1-24/Domain: signal sequence #status predicted
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Residues: 1-116 <FUH>
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A;Cross-references: GB:V00640; GB:J00946; NID:g64028; PIDN:CAA23986.1; PID:g64029 R;Goodman, R.H.; Jacobs, J.W.; Chin, W.W.; Lund, P.K.; Dee, P.C.; Habener, J.F. Proc. Natl. Acad. Sci. U.S.A. 77, 5869-5873, 1980 A;Title: Nucleotide sequence of a cloned structural gene coding for a precursor of A;Reference number: A93860; MUID:81077276; PMID:6108560
                                                                                                                                                                                                                                                                                               A; Reference number: A91087; A; Accession: A91087
                                                                                                                                                                                                                                                                                                                      A; Title: Isolation and characterization of A; Reference number: A91087; MUID:80046482;
                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Experimental source: islet tissue (endocrine p R; Goodman, R.H.; Jacobs, J.W.; Chin, W.W.; Lund, Proc. Natl. Acad. Sci. U.S.A. 79, 1682, 1982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C; Species: Lophius americanus (American goosefish) C; Date: 31-Mar-1981 #sequence_revision 31-Mar-1981 C; Accession: A93236; A93860; A91087; A01433
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C; Species: L
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                                                                                                                                                                                                                                                          A; Molecule type: protein A; Residues: 108-121 < NOE>
                                                                                                                                                                                                                                                                                                                                                           R; Noe, B.D.; Spiess, J.; Rivier, J
Endocrinology 105, 1410-1415, 1979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 2-20,'V',22-82,'E',84-121 <GOO>
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A; Residues: 1-121 < HO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Title: Cloning and sequence analysis of cDNAs encoding A; Reference number: A93236; MUID:81052423; PMID:6107860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-116 <NAT>
                                                                                                                                                           F;110-121/Disulfide
                                                                                                                                                                                                                                                                                                                                                                                              A; Contents: annotation; erratum
                                                                                                                                                                                                                                                                                                                                                                                                                     A; Reference number: A93905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Accession: A93860
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                                                                                                 Best
                                                                              Matches
                                                                                                                 Query Match
                                                                                                                                                                                            1-24/Domain: signal sequence
                                                                                                                                                                                                                     Keywords: neuropeptide
                                                                                                                                                                                                                                         Superfamily: somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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12 LLVLLISLTASISCSFAG---QRDSKLRLLLHRYPLQGSKQDMTRSALAELLLSDLLQGE
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                                       LLLLLSGATATAALPLEGGPTGRDSE----HMQEAAG---
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                                                                                             16.5%;
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                                                                                               Score 91.5;
Pred. No. 0
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PMID:387385
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d, P.K.; Dee,
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51

-- AWWFEWTSQASAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSC

of 1

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N;Contains: somatostatin 14 (SS-14); somatostatin 28 (SS-28) C;Species: Homo sapiens (man) C;Date: 17-Dec-1982 #sequence_revision 17-Dec-1982 #text_change C;Accession: A43614; A01430; S09381; S50024 R;Shen, L.P.; Rutter, W.J. Science 224, 168-171, 1984 Science 224, 168-171, 1984 A;Ritle: Sequence of the human somatostatin I gene. A;Reference number: A43614; MUID:84146798; PMID:6142531
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                                                                                                                                                                                                                                                                                                                                                                                                             A; Description: inhibits the secretion of a number of peptide C; Superfamily: somatostatin C; Keywords: hormone; hypothalamus; neuropeptide C; Keywords: hormone; hypothalamus; neuropeptide SIG>
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EMBO J. 8,
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A; Residues: 1-116 <SHE>
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A; Residues: 1-116 <SH2>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Function:
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                                                                                                                                                                                                                                                Local
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. J. 303, 263-268, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human seminal plasma contains somatostatin-64.
nce number: S50024; MUID:95031969; PMID:7945250
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8, 2911-2916, 1989
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                                                                              IGEEAREVARRQE---
                               EPEDLSQAAEQDEMRLELQRSANSNPAMAPRERKAGCKNFFWKTFTSC
                                                                                                                             ALAALSIVLALGCVTGAPSDPRLRQFLQKSLAAAAGKQELAKYFLAELLSEPNQTENDAL
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inhibits
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29.6%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Title: Prosomatostatin 1-64 is a major product of somatostatin A; Reference number: A34109; MUID:89278131; PMID:2567292 A; Accession: A34109
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                                              A; Experimental source: intestine R; Schally, A.V.; Huang, W.Y.; Chang, R.C.C.; Arimura, Proc. Natl. Acad. Sci. U.S.A. 77, 4489-4493, 1980
                                                                                                                        A; Molecule type: protein A; Residues: 65-92 < PRA>
                                                                                                                                                                    A; Reference number: A91273; A; Accession: A91273
                                                                                                                                                                                                                     A; Title: N-terminally extended somatostatin:
                                                                                                                                                                                                                                      R; Pradayrol, L.; Jornvall, FEBS Lett. 109, 55-58, 1980
                                                                                                                                                                                                                                                                                          A; Molecule type: protein A; Residues: 1-32 <SC3>
                                                                                                                                                                                                                                                                                                                                        A; Reference number: A24222;
A; Accession: A24222
                                                                                                                                                                                                                                                                                                                                                            A; Title: Isolation and characterization of proSS1-32, a A; Reference number: A24222; MUID:86030691; PMID:2865169
                                                                                                                                                                                                                                                                                                                                                                                                        R; Schmidt, W:E.; Mutt, V.; FEBS Lett. 192, 141-146, 1
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C:Superfamily: somatostatin
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A; Residues: 1-116 <TRA>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         somatostatin I precursor - pig (fragment)
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Sutcliffe, J.G.
16.96-1700, 1988
Id. Sci. U.S.A. 85, 1696-1700, 1988
Id. emulsion-enhanced DNA-driven subtractive
Phar: A28968; MUID:88144503; PMID:2894033
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29.6%;
                                                                                                                                                                                               MUID:80113258;
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Pred. No.
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PMID:7353633
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                                                                       Redding,
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A;Residues: 1-115 <TOS>
A;Cross-references: GB:U68136; NID:g1890650; PIDN:AAC60093.1; PID:g1890651 C;Comment: This protein acts both as a neurotransmitter/neuromodulator and a C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N;Alternate names: PSSI protein
C;Species: Rana ridibunda (laughing frog)
C;Date: 11-Apr-1997 #sequence_revision 09-May-1997 #text_change 16-Jul-1999
C;Accession: JC6166
R;Tostivint, H.; Lihrmann, I.; Bucharles, C.; Vieau, D.; Coulouarn, Y.; Fournier, Proc. Natl. Acad. Sci. U.S.A. 93, 12605-12610, 1996
A;Title: Occurrence of two somatostatin variants in the frog brain: Characterization R;Reference number: JC6166; MUID:97057290; PMID:8901629
N;Contents: brain
comatostatin-related protein - Atlantic hagfish
C;Species: Myxine glutinosa (Atlantic hagfish)
C;Date: 21-May.1990 #sequence_revision 21-May-1990 #text_change
C;Accession: A32271
                                                                                         RESULT
A32271
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C;Superfamily: somatostatin
C;Keywords: brain; hormone
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R;Bersani, M.; Johnsen, A.H.; Holst, J.J.
FEBS Lett. 279, 237-239, 1991
A;Title: Oxidation/reduction explains heterogeneity of pancreatic somatostatin.
A;Title: Oxidation/reduction explains heterogeneity of pancreatic somatostatin.
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A;Acession: A90398
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A; Residues: 65-92 <SCH>
A; Experimental source: hypothalamus
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;Reywords: hormone; hypothalamus; intestine; neuropeptide
;1-64/Domain: propeptide #status experimental <PRO>
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 87; DB 1;
Pred. No. 0.045;
7; Mismatches 1
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Pred. No.
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0.071;
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                       31-Dec-1993
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Дb Qγ 밁 20

preprosomatostatin

SS-14 -

channel catfish

RESULT

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A; Molecule type: mRNA
A; Residues: 82-108 <TAY>
A; Residues: 82-108 <TAY>
A; Cross references: GB.J00944
A; Cross references: GB.J00944
R; Andrews, P.C.; Dixon, J.E.
J. Biol. Chem. 256, 8267-8270, 1981
J. Biol. Chem. 150Jation and structure of a peptide hormone predicted from a mRNA sequence.
A; Reference number: A92334; MUID:81264223; PMID:6114953
A; Accession: A92334; MUID:81264223; PMID:6114953
                                                                                                                                                                                                                                                                                                                                     F;1-24/Domain: signal sequence #status predicted <SIG>
F;25-100/Domain: propeptide #status predicted <PRO>
F;101-114/Product: somatostatin-14 #status experimental <MAT>
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C;Date: 30-Jun-1980 #sequence_revision 31-Dec-1993 #text_change 18-Jun-1999
C;Accession: S00292; A93897; A92334; A01435
R;Minth, C.D.; Taylor, W.L.; Magazin, M.; Tavianini, M.A.; Collier, K.; Weith,
J. Biol. Chem. 257, 10372-10377, 1982
A;Title: The structure of cloned DNA complementary to catfish pancreatic somatc A;Reference number: S00292; MUID:82265698; PMID:6179939
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Endocrinology 122, 1855-1859, 1988
A;Title: Primary structures of somatostatins from the islet organ of the hagfish sugg.
A;Reference number: A32271; MUID:88195948; PMID:2896118
A;Accession: A32271
                                                                                                                                                                                                                                                                                                         F;101-114/Product: somatostatin-14 *status expeF;103-114/Disulfide bonds: *status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                    C;Superfamily: somatostat
C;Keywords: neuropeptide
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A; Residues: 1-114 < MIN>
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                                                 PLIGEEAREVA-----RRQEGAPPQQSARRDRMPCRNFFWKTFSSC 104
  VLDSDEVSRAAESEGARLEMERAAGPMLAPRERKAGCKNFFWKTFTSC
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C;Species: Ictalurus punctatus (channel catfish)
C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 16-Jul-1999
C;Accession: I50798
R;Dixon, J.E.; Andrews, P.C.
Adv. Exp. Med. Biol. 188, 19-29, 1985
A;Title: Somatostatins of the channel catfish.
A;Reference number: I50798; MUID:85303576; PMID:2863931
A;Accession: I50798
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-114 CDIX>
A;Cross-references: GB:M25903; NID:g213339; PIDN:AAA49339.1; PID:g213340
C;Superfamily: somatostatin
                                                                                                               Query Match 15.0%; Score 83; DB 2; Length 114; Best Local Similarity 37.0%; Pred. No. 0.14; Matches 17; Conservative 7; Mismatches 12; Indels
10;
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Search completed: June 23, 2003, 15:57:30 Job time: 42 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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197.955 Million cell updates/sec
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554
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  SwissProt_40:*
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P87385 rana ridibu
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Result No.

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Minimum DB Maximum DB

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Scoring table: Sequence:

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on:

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		Q92622 homo sapien								Q9ygh4 carassius a	Q62925 rattus norv

ALIGNMENTS

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EMBL; ABO00263; BAA19770.1; - GENBL; AF013252; AAB66895.1; - GENBW; HGNC:2257; CORT. MIM; 602784; - InterPro; IPR004250; Somatost Pfam; PF03002; Somatostatin; Cleavage on pair of basic res SIGNAL 1 18 PEPTIDE 77 105 PEPTIDE 89 105	SWING EURON BY Fied fied ties	-!- SUBCELLULAR LO -!- TISSUE SPECIFI THE CORTEX AND -!- SIMILARITY: BE	Genomics 42:499-506(1997)I- FUNCTION: BINDS TO ALL SUBTYPES. IT ALSO INHI	SEQUENCE FROM N.A. SEQUENCE FROM N.A. MEDLINE-97349120; PubMed-9205124; de Lecea L., Ruiz-Lozano P., Danielson feye P.E., Frankel W.N., Sutcliffe J.G "Cloning, mRNA expression, and chromos	TISSUE-Brain; MEDLINE-97236300; Frukusumi S., Kita- Miyamoto M., Hinu "Identification a: Cortistatin-like Biochem. Biophys.	1 11 7 16	15-JUL-1998 15-JUL-1998 15-JUL-1998 16-OCT-2001 Cortistatin	DRT_HUMAN CORT_HUMAN
63; BAA19 52; AAB66 2257; COF R004250; 2; Somatc pair of k 1 77 10 89	SS-PROT entry is conthe Swiss Institution of the Swiss Institution on profit institution on profit institution of this statement requires a license an email to license	ILAR LOCATION: Secre SPECIFICITY: EXPRESS EX AND HIPPOCAMPUS. TY: BELONGS TO THE	499-506(199 1: BINDS TO 3. IT ALSO I	FROM N.A. 97349120; PubMed=9 L., Ruiz-Lozano P ., Frankel W.N., S ., mRNA expression,	room, N.A. ain; 7236300; PubMed=91 8., Kitada C., Tak M., Hinuma S., Kit cation and charact in-like peptide."; Biophys. Res. Comm	man) (oa; (ia;	(Rel. 36, 0 (Rel. 36, I (Rel. 40, I precursor	STANDARD;
	ROT entry is copyright. It is programs is swiss Institute of Bioinformatics Bioinformatics Institute. There Profit institutions as long as this statement is not removed. Unlike a license agreement (See httmail to license@isb-sib.ch).	SUBCELLULAR LOCATION: Secreted. TISSUE SPECIFICITY: EXPRESSED IN A SU THE CORTEX AND HIPPOCAMPUS. SIMILARITY: BELONGS TO THE SOMATOSTAT	9-21. 9-21. 9-206(1997). BINDS TO ALL HUMAN SOIT ALSO INHIBITS CAMP	PubMed=9205124; -Lozano P., Danielson F L W.N., Sutcliffe J.G.; pression, and chromosom	2512 ekaw ano eriz	Chordata; Cran. Primates; Cata	Created) Last sequence up Last annotation [Contains: Cort.]); PRT;
Hormone; Signal. IAL. TATIN-29 (POTENTIA	It is produced through a colinformatics and the EMBL of the There are no restrictions long as its content is removed. Usage by and for nt (See http://www.isb-sib.cl.ch).	SUBSELLULAR LOCATION: Secreted. TISSUE SPECIFICITY: EXPRESSED IN A SUBSET OF GABAERGIC CELLS IN THE CORTEX AND HIPPOCAMPUS. SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.	99-506(1997). BINDS TO ALL HUMAN SOMATOSTATIN RECEPTOR (SSTR) IT ALSO INHIBITS CAMP PRODUCTION INDUCED BY FORSKOLIN	P.E., Peelle-Kirley J., .; omal mapping of mouse and	2; R., Kizawa H., Sakamoto J., K., Fujino M.; ation of a novel human 232:157-163(1997).	Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.	Created) Last sequence update) Last annotation update) [Contains: Cortistatin-29; Cortistatin-17]	105 AA.
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                                                                                                                                                                                                                                                                                              use by non-profit institutions as long modified and this statement is not removed. entitles requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     properties.";
Nature 381:242-245(1996).
-i- FUNCTION: NEUROPEPTIDE WITH NEURONAL DEPRESSANT
                                                                                                                                                                                                                     Pfam; PF03002;
                                                                                                                                                                                                                                                   EMBL; U51919; AAC52585.1;
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                                                                                                                                                                                                                                                                                                                                               the European Bioinformatics Institute.
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MEDLINE=96208649; PubMed=8622767;
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lecea L., Criado J.R., Prospero-Garcia O., hweitzer P., Danielson P.E., Dunlop C.L.M.,
                                                                                                                                                                                                                                                                                                                                                                                                           AREAS AND IN THE DENTATE GYRUS. A TRANSIENT EXPRESSION IS OBSERVED IN THE HILAR REGION AT P16.
SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MODULATING PROPERTIES.
SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: INTERNEURONS IN THE CEREBRAL CORTEX AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYPPOCAMPUS.

DEVELOPMENTAL STAGE: THERE IS A TRANSIENT INCREASE IN CORTISTATINEXPERSING CELLS IN THE SECOND POSTNATAL WEEK IN ALL CORTICAL
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                                                   l Similarity
58; Conserv
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105 /
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(Rel. 35, Last sequence update)
(Rel. 40, Last annotation update)
precursor [Contains: Cortistatin-29;
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                                                       Conservative
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                                                                                                                                                  basic residues; Hormone;
27 POTENTIAL.
112 CORTISTATIN-29
112 CORTISTATIN-14.
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Rodentia;
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                                                                                                                      12201 MW;
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                                                                      53.2%;
55.8%;
                                                                                                                                                                                                                                  Somatostatin.
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                   SIMILARITY. FF1FBFC76CBD7991 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 SIMILARITY.
09578F4520201551 CRC64;
                                                       Mismatches
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No. 3
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                                                                       .7e-24;
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                                                                                                                                                                   (POTENTIAL)
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Siggins
                                                                                 Length 112;
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ID CORT_M
AC P56469
AC P56469
DT 15-JUL
DT 16-OCT
DE CORTIS
GN CORT.
OS MUS MU
OC MAMMAIN
OX NCBI_T
RN [1]
RP SEQUEN
RX MEDLIN
RX MEDLIN
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PLSPGLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQAS

6

Last sequence update)
Last annotation update)

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Query Match
Best Local
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SMS2_RANRI
P87385;
16-OCT-2001
16-OCT-2001
16-OCT-2001
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JUL-1998
15-JUL-1998
16-OCT-2001
                                                                            RANRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sorsend an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                 Cleavage on pair of basic residues; Hormone; Signal SIGNAL 1 25 POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          THE CORTEX AND HIPPOCAMPUS.
-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     preprocortistatin.";
Genomics 42:499-506(1997).
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: EXPRESSED I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=97349120; PubMed=9205124;
de Lecea L., Ruiz-Lozano P., Danielson
Foye P.E., Frankel W.N., Sutcliffe J.G.
"Cloning, mRNA expression, and chromosc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. STRAIN-C57BL/6J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; AF013253; AAB66896.1; ~.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus (Mouse)
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                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                PLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQAS
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                                                                                                                                    SSTPVGGGTPGLSKSQERPPPQQPPHLDKKPCKNFFWKTFSSCK
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97
109
 (Rel.
(Rel.
(Rel.
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(Rel. 36, Last sequence update)
(Rel. 40, Last annotation update)
precursor [Contains: Cortistatin-14].
                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                     AA;
                                                            STANDARD;
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11613 MW;
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Rodentia;
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                               Created)
                                                                                                                                                                                                                                                                         Score 272;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                     CORTISTATIN-14.
BY SIMILARITY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chromosomal mapping
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                                                            103
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Matches
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                                                                                                             SMSB_CARAU
Q9YGH3;
16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fournier A., Conlon J.M., rausa ... Procedurence of two somatostatin variants in the frogramaterization of the cDNAs, distribution of the receptor-binding affinities of the peptides.";

Broc. Natl. Acad. Sci. U.S.A. 93:12605-12610(1996).
                                                                                                                                                                                                                                                                                                                                                                                DISULFID
SEQUENCE
          Carassius auratus (Goldfish).
Eukaryota; Metazoa; Chordata;
Actinopterygii; Neopterygii;
                                                                  16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation updat
Somatostatin IB precursor [Contains: [Pro12]
                                                                                                                                                       CARAU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vaudry H., Chartrel N., Conlon J.M.; "Isolation of [Pro2,Met13]somatostatin-14 and somatostatin-14 from the frog brain reveals the existence of a somatostatin gene family in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Somatostatin 2 precursor (PSS2) [Contains: (S-2) (SS2)].
Rana ridibunda (Laughing frog) (Marsh frog)
                                                                                                                                                                                                                                                                                                                                                                                                                                       SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cleavage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Biochem. Biophys. Res. Commun. 188:477-482(1992).
-I- FUNCTION: SOMATCSTATIN INHEBITS THE RELEASE OF .
-I- SUBCELLULAR LOCATION: Secreted.
-I- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=93038702; PubMed=1358069;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amphibia; Batrachia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
                                                        [Pro2]Somatostatin-14]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE OF 90-103.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-97057290; PubMed-8901629;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                         Local
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35; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                    9
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                                                                                                                                                                                                                                      GEEAREVARRQEGAP------PQQSARRDRMPCRNFFWKTFSSC
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                                                                                                                                                                                                                                                                                                                                                                                 103
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                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conlon J.M., Vaudry H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 precursor (PSS2) [Contains: [Pro2, Met13] somatostatin-14
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                                                                                                                                                                                                                                                                                                                                                                                AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Somatostatin;
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21
87
103
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11647
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Somatostatin
                                                                                                                                                                                                                                                                                                                                      20.3%;
                                                                                                                                                                                                                                                                                                                                                                                MW;
                                                                                                                                                                                                                                                                                                                                                                                                      residues; Hormone; Multigene fa
BY SIMILARITY.
BY SIMILARITY.
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                          13;
                                                                                                                                                                                                                                                                                                                         Score 112.5;
Pred. No. 4.2e
13; Mismatches
             ; Craniata;
Teleostei;
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                                                                                                                                        PRT;
                                                                                                                                                                                                                                                                                                                                                                             ADBA7329857EF77A CRC64;
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                                                                                                                                        111
                                                                    [Pro12]somatostatin-24;
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           Vertebrata; Euteleostomi;
Ostariophysi; Cypriniform
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hes 37;
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                                                                                                                                                                                                                                                                                                                                                 Length 103;
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Ranidae; Rana
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Best Local
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01-AUG-1992
16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SMS_BOVIN P26917;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BOVIN
  Furu L.M., Kazmer G.W., Strausbaugh L., "Cloning and characterization of the boy J. Anim. Sci. 77:492-493(1999).
-I- FUNCTION: SOMATOSTATIN TNHTRITES TUP
                                                                                                                                                                                                                                  MEDLINE-88288237; Pul
Su C.J., White J.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                    SST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                   Chan
                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                       Bovidae; Bovinae;
NCBI_TaxID=9913;
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                                                                                    MEDLINE=99198780; PubMed=10100681;
                                                                                                        STRAIN-Holstein;
                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                              Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Somatostatin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Molecular cloning
                                                                                                                                                                      "Structure and evolution of somatostatin Mol. Endocrinol. 2:209-216(1988).
                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DISULFID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       goldfish.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam; PF03002; Somatostatin;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- SUBCELLULAR LOCATION: Secreted.-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE-Brain,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=7957;
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                                                                                                                                                                                                                                                                                                                                                                                                         (Bovine)
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(Rel. 23, Last sequence update)
(Rel. 40, Last annotation updat
n precursor [Contains: Somatosta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             111 AA;
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W., Li W.H., Luo
                                                                                                                                                                                                                                                                                                                                               Bos.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   [PRO12]SOMATOSTATIN-24
[PRO2]SOMATOSTATIN-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 112.5;
Pred. No. 4.
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Mismatches 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   tion update)
Somatostatin-28;
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                                          bovine somatostatin gene.";
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  RELEASE OF
                                                                                                                                                                                                                                     Frazier
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Best Local S
Matches 34
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                                   the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license.agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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DISULFID
SEQUENCE
                                                                                                                                                                                                                                STRAIN-Ile de France;
MEDLINE-99094691; PubMed-9880082;
Bruneau G., Tillet Y.;
                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea; Bovidae; Caprinae; Ovis.
EMBL;
                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation
                                                                                                                                                                          in the ewe hypothalamus.";
Peptides 19:1749-1758(1998).
-!- FUNCTION: SOMATOSTATIN INHIBITS THE
                                                                                                                                                                                                                                                                                              NCBI_TaxID=9940;
                                                                                                                                                                                                                                                                                                                                                   Ovis aries (Sheep).
                                                                                                                                                                                                                                                                                                                                                                                Somatostatin precursor (Contains: Somatostatin-28;
                                                                                                                                                                                                                                                                                                                                                                                       046688;
15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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EMBL; U97077;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long modified and this statement is not removed entitles requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                                                                                           "Localization of the preprosomatostatin-mRNA by in the ewe hypothalamis.":
                                                                                                                                                                                                                                                                         SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SIGNAL
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SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
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Y15267;
           AF031488; AAC04697.1;
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BELONGS TO THE SOMATOSTATIN FAMILY.
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12688 MW;
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BY SIMILARITY.
SOMATOSTATIN-28.
SOMATOSTATIN-14.
BY SIMILARITY.
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                                                                                                                                                                            RELEASE
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RESULT 8
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PEPTIDE
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                                                                                                                        Ptam; PF03002; Somatostatin;
                                                                                                                                              EMBL; L42325; AAA67099.1;
                                                                                                                                                                entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation
                                                                  PEPTIDE
                                                                                                                                                                                                                                                      -!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                        Regul. Pept. 67:145-152(1996).
                                                                                                                                                                                                                                                                                                                                            Dickinson C.J.,
                                                                                                                                                                                                                                                                                                                                                      MEDLINE=97142297; PubMed=8988514;
                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
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01-FEB-1996 (Rel.
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33, Last sequence update)
40, Last annotation update)
ursor [Contains: Somatostatin-28;
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28.4%;
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BY SIMILARITY.
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residues; Hormone;
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SOMATOSTATIN-28.
SOMATOSTATIN-14.
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Pred. No.
          Score 95.5;
Pred. No. 0
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                                                                 SOMATOSTATIN-14.
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P01167;
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PEPTIDE
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                                                                                                                                                                                                                                                                                                                                                                                                                      use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an
                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -1- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF -1- SUBCELLULAR LOCATION: Secreted. -1- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Protopterus annectens (African lungfish).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Dipnoi; Lepidosireniformes; Protopteridae; Protopterus.
NCBI_TaxID=7888;
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MEDLINE=99326690; PubMed=10398054;
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16-0CT-2001
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                                                                                                                                                                                                                     Local
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                                         STANDARD;
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precursor (PSS2) [Contains:
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Pred. No.
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                                        PRT;
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  Fuhrmann G., Heilig R., Kempf J., Ebel A.;
"Nucleotide sequence of the mouse preprosomatostatin gene.";
Nucleic Acids Res. 18:1287-1287(1990).
-!- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTRO!
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                     SPECIES=Mouse; TISSUE=Brain; MEDLINE=90206793; PubMed=1969620;
                                                                                                                        "A new prosomatostatin-derived peptide reveals a prohormone cleavage at monobasic sites."; Science 238:1126-1129(1987).
                                                                                                                                                                          SPECIES=Rat; STRAIN=Sprague-Dawley; MEDLINE=88070564; PubMed=2891188;
                                                                                                                                                                                                                         medullary thyroid carcinoma.";
J. Biol. Chem. 257:1156-1159(1982).
                                                                                                                                                                                                                                                                                                                    Proc.
                                                                                                                                                                                                                                                                                                                         MEDILINE=84221954; PubMed=6145156;
MEDILINE=84221954; PubMed=6145156;
Montminy M.R., Goodman R.H., Horovitch S.J., Habener J.F.;
"Primary structure of the gene encoding rat preprosomatostatin.";
Proc. Natl. Acad. Sci. U.S.A. 81:3337-3340(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-85303584; PubMed=2863939; Goodman R.H., Montminy M.R., Low M.J., "Biosynthesis of rat preprosomatostatir "Biosynthesis of tat [Page 188]. Adv. Exp. Med. Biol. 188:31-47(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=83238516; PubMed=6134734; Argos P., Taylor W.L., Minth C.D., I "Nucleotide and amino acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Somatostatin precursor Somatostatin-14]. SST OR SMST.
                                                                                                                                                                                                    SEQUENCE OF 25-34
                                                                                                                                                                                                                                          "Somatostatin-28 encoded in a cloned
                                                                                                                                                                                                                                                                            MEDLINE=82120034; PubMed=6120163;
                                                                                                                                                                                                                                                                                           SPECIES-Rat;
                                                                                                                                                                                                                                                                                                                                                                                  SPECIES-Rat
                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=83213516; PubMed=6133871; Goodman R.H., Aron D.C., Roos B.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           preprosomatostatins.";
J. Biol. Chem. 258:8788-8793(1983).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SPECIES-Rat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-OCT-2001 (Rel. 40, Last annotation update)
Somatostatin precursor [Contains: Antrin; Somatostatin-28,
                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                              Benoit R., Ling N., Esch F.
                                                                                                                                                                                                                                                                                                      SEQUENCE OF 38-116 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
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membranes.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        258:5570-5573(1983).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; PubMed=6148343;
Hayes T.E., Magazin M.D., Minth C.
                                                                                                                                                                                                                                                                                                                                                                                                                   preprosomatostatin."; 188:31-47(1985).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dixon
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                                                                                                                                                                                                                                                     Habener J.F.;
cDNA obtained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                            Habener
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                processing
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Matches 35
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EMBL; V01271; CAA24579 1; -
EMBL; J00787; AAA42164 1; -
EMBL; M25890; AAA42167 1; -
EMBL; J00788; AAA42162 1; -
                                       This SWIS
between
the Europ
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01-OCT-1993
01-OCT-1993
16-OCT-2001
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                                                                                                                                                      YONEKURA H., Okamoto H.;
Submitted (JUN-1991) to the EMBL/GenBank/DDBJ
Submitted (JUN-1991) THE RELEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PEPTIDE
PEPTIDE
    modified and
                    use
                                                                                                                                                                                                                          Nata
                                                                                                                                                                                                                                             TISSUE=Pancreas;
                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                                                                                            Archosauria;
                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                   Gallus gallus (Chicken)
                                                                                                                                                                                                                                                                                                                                                                                                                      Somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SMS_CHICK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This
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InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DISULFID
                  by non-profit institutions as long
                                                                                                                     SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                          SUBCELLULAR LOCATION: Secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             European Bioinformatics Institute.
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                                                       SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SWISS-PROT entry is copyright. It is produced through en the Swiss Institute of Bioinformatics and the EN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                      Kobayashi T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LLLLLLSGATATAALPLE-GGPTGRDS-----EHMQE--AAGIRKSSLLT-FLAWWFEWT
  non-profit and this st
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(Rel.
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Last annotation update)
r [Contains: Somatostatin-28;
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                                                                                                                                                                                                                    Karahashi K.,
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14; Mismatches
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SOMATOSTATIN-14
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; Galliformes; Phasianidae; Phasianinae;
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-> H (IN REF. 6).
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                                                                                                                                                                                                                      Kato S.,
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                There are no restrictions ong as its content is in
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Usage
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Best Local
                                                                                                                                                                                                                                                                         somatostatin precursors f
anglerfish.";
Nature 288:137-141(1980).
                     Endocrinology 105:1410-1415(1979).
-!- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF
-!- SUBCELLULAR LOCATION: Secreted.
                                                      Noe B.D., Spiess J., Rivier J.E "Isolation and characterization pancreatic islet.";
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                                                                                                                                      Goodman R.H., Habener J.F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SMS1
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PROPEP
                                                                                         SEQUENCE OF 108-121.
MEDLINE=80046482; Pu
                                                                                                                             Proc.
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                                                                                                                                                                                   Proc.
                                                                                                                                                                                              precursor of pancreatic
                                                                                                                                                                                                                     Goodman R.H.,
Habener J.F.;
                                                                                                                                                                                                                                          MEDLINE=81077276;
                                                                                                                                                                                                                                                      SEQUENCE OF 2-121
                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A. MEDLINE=81052423;
                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                             Lophius americanus (American goosefish) (Anglerfish).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; -
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Paracanthopterygii; Lophiiformes; Lophiidae; Lophius.
                                                                                                                                                                                                                                                                                                                                                                                                                           Somatostatin I precursor [Contains:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                 "Nucleotide sequence of a cloned structural gene precursor of pancreatic somatostatin."; Proc. Natl. Acad. Sci. U.S.A. 77:5869-5873(1980)
                                                                                                                                                                                                                                                                                                                      Hobart P.M.,
                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=8073;
                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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          SIMILARITY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                66
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                                                                                                                                                                                                                                                                                                            Crawford R., Shen L., Pictet R., sequence analysis of cDNAs encod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25
89
                                                                                                                                                                                                                             -121 FROM N.A.
276; PubMed=6108560;
Jacobs J.W., Chin W.W.,
                                                                                                                                                 Jacobs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
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          BELONGS
                                                                             PubMed=387385;
J., Rivier J.E.,
                                                                                                                             Sci.
                                                                                                                                                                                                                                                                                                                                  PubMed=6107860;
                                                                                                                                                 J.W.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16.7%;
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         TO THE SOMATOSTATIN FAMILY
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                                                                                                                                                 Chin W.W.,
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BY SIMILARITY.
SOMATOSTATIV. 28.
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Pred. No. 0
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                                                                                                                             79:1682-1682(1982)
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RESULT 13
SMSA_CARAU
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Best Local S
Matches 35
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SEQUENCE
                                                                                                                                                                 "Cloning and characterization of cDNAs encoding preprosomatostatin-I and -II from goldfish brain.";
Submitted (NOV-1995) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: SOMATOSTATIN INHIBITS THE RELEASE OF SOMATOTROPIN.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                               16-0CT-2001
16-0CT-2001
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                                                                                                                                                                                                                                                                                            Cyprinidae; Car
NCBI_TaxID=7957
                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata Actinopterygii; Neopterygii; Cyprinidae; Carassius.
       Cleavage on
                                 EMBL; U40754; AAD09359.1; -. InterPro; IPR004250; Somatostatin.
                                                                                                                     the
                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                        TISSUE-Brain;
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16-OCT-2001
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                    Pfam; PF03002; Somatostatin;
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                                                                                                                                                                                                                                             Lin X.-W.,
                                                                                                       European Bioinformatics Institute. The by non-profit institutions as long
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pterygii; Teleostei;
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          Hormone;
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Somatostatin-26;
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RESULT 14
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                                                        This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sentites.com/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/markets/server/mark
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"Human somatostatin I: sequence of the cDNA.";
Proc. Natl. Acad. Sci. U.S.A. 79:4575-4579(1982).
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                                                                                                                                                                                                                                                                                                                                        <del>: :</del>
                                                                                                                                                                                                                                                                                                                                                                              "Phenol emulsion-enhanced DNA-driven subtractive cDNA cloning: isolation of low-abundance monkey cortex-specific mRNAs."; Proc. Natl. Acad. Sci. U.S.A. 85:1696-1700(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-88144503; PubMed=2894033; Travis G.H., Sutcliffe J.G.;
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"Sequence of the human somatostatin I
Science 224:168-171(1984).
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16-OCT-2001
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                                                                                                                                                                                                                    this is a synthetic cyclic analog known as octreotide or : 201-995. Used for the treatment of a variety of disorders including acromegaly and the symptomatic treatment of card tumors and vasoactive intestinal peptide tumors. SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
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r [Contains: Somatostatin-28;
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                                                                                                                                                                                                                                                                    carcinoid
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Matches 32
TISSUE-Hypothalamus;
MEDLINE-81054799; PubMed-6107906;
Schally A.V., Huang W.-Y., Chang R.C.C., Arimura A.,
Millar R.P., Hunkapiller M.W., Hood L.E.;
"Isolation and structure of pro-somatostatin: a putat
precursor from pig hypothalamus.";
                                                                                                        TISSUE-Intestine;
TISSUE-Intestine;
MEDLINE-80113256; PubMed=7353633;
Pradayrol L., Joernvall H., Mutt V.,
"N-terminally extended somatostatin:
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P01168;
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MEDLINE-86030691; PubMed=2865169;
Schmidt W.E., Mutt V., Kratzin H.,
                                                                                                                                                                                                                                                                                    expression in pancreas and gut.";
J. Biol. Chem. 264:10633-10636(1989).
                                                                              SEQUENCE OF 65-92
                                                                                                                                                                       SEQUENCE OF 65-92.
                                                                                                                                                                                          "Isolation and characterization of proSS1-32, a peptide derived the N-terminal region of porcine preprosomatostatin."; FEBS Lett. 192:141-146(1985).
                                                                                                                                                                                                                                                                                                 Bersani M., Thim L., Baldissera r.o.n., a
"Prosomatostatin 1-64 is a major product
                                                                                                                                                                                                                                                                                                                               SEQUENCE OF 1-64.
MEDLINE=89278131;
                                                                                                                                                                                                                                                                                                                                                                             Sus scrofa (Pig).
Eukaryota; Metazoa;
Mammalia; Eutheria;
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01-OCT-1989 (Rel.
16-OCT-2001 (Rel.
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PIR; A28968; A28968.
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                                                                                                    Lett. 109:55-58(1980).
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Pred. No.
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MEDLINE-76136331; PubMed=1252409;

MEDLINE-7636331; PubMed=1252409;

Schally A.V., Dupont A., Arimura A., Redding T.W., Nishi N.,

Schally A.V., Dipont A., Arimura A., Redding T.W., Nishi N.,

Linthicum G.L., Schlesinger D.H.;

"Isolation and structure of somatostatin from porcine hypothalami.";

Biochemistry 15:509-514(1976).
                                                                                                                                                       SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                     use by non-profit institutions as long modified and this statement is not removed
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                                                                                                                                                                                                                                Cleavage
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-1- SIMILARITY: BELONGS TO THE SOMATOSTATIN FAMILY.
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                                                                                                      1 Similarity
15; Conser
                                                   DEMRLELQRSANSNPAMAPRERKAGCKNFFWKTFTSC 92
                                                                 EEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSC 104
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SOMATOSTATIN-14.
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O90xe0 acipenser t
O90xe1 acipenser t
O8xem7 brucella me
O8zb18 yersinia pe
O91xn4 arabidopsis
O8xu92 ralstonia s
O92m60 rhizobium m
                                  Q90y39 catostomus
Q90y41 gnathonemus
Q90y40 chitala chi
Q90gdd3 oryza sativ
                       Q9sdd3 ory:
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Q99mr0 mus
075421 homo
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5		⇉	P89445 herpes simp			Q9krq4 vibrio chol			Q.			ω			9			a	ω			ω	Q8rqp8 streptomyce		Q98f77 rhizobium l	Q95yu0 leishmania		Q8wzq0 neurospora

ALIGNMENTS

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RESULT 1
Q9R1P8
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                                     Query Match
Best Local :
      Matches
                                                                                                                                                                               Calbet Murtro M., de Lecea L.;
Submitted (FEB-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF050156; AAD51127.1; -.
MGD; MGI:109538; Cort.
InterPro; IPR004250; Somatostatin.
Pfam; PF03002; Somatostatin; 1.
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O9R1P8;

O1-MAY-2000 (TrEMBLrel. 13,

O1-MAY-2000 (TrEMBLrel. 13,

O1-DEC-2001 (TrEMBLrel. 19,
                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. STRAIN-129/SV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Calbet M., Guadano-Ferraz A., Spier A.D., Maj M., Sutcliffe J.G., Przewlocki R., de Lecea L.; "Corristatin and somatostatin mRNAs are differentially regulated in response to kainate.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A. STRAIN=129/SV;
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Mammalia; Eutheria;
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   53;
                                     Similarity
                                                                                                                                     105 AA; 11061 MW;
      Conservative
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Rodentia;
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51.0%;
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                    Score 248; Db 1.
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      Mismatches
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Matches 32
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Eukaryota; Metazoa; Chordata; Craniata;
Actinopterygii; Chondrostei; Acipenserifi
                                                                                                                                                                                                                                                                                          01-DEC-2001 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
                               encoding two somatostatin precursors.";
J. Comp. Neurol. 0:0-0(2001).
EMBL; AF395849; AAL13248.1; -.
                                                                              "Polygenic expression of somatostatin in transmontanus: molecular cloning and dist
                                                                                                              Trabucchi M., Tostivint H., Lihrmann I.,
Dores R.M., Vaudry H.;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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01-DEC-2001 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
                                                                                                                                            TISSUE-BRAIN;
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Dores R.M., Vaudry H.;
"Polygenic expression of somatostatin in
transmontanus: molecular cloning and dist
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                  InterPro;
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PF03002;
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                  IPR004250;
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formes; Acipenseridae;
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NCBI_TaxID=632;
                rersinia
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01-MAR-2002
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01-MAR-2002 (TrEMBLrel. 20, 01-MAR-2002 (TrEMBLrel. 20, 01-JUN-2002 (TrEMBLrel. 21,
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EMBL; AE009501; AAL52841.1; -
EMBL; AE009531; AAL52084.1; -
Interpro; IPR002086; Aldehyde_dehydr.
PROSTTE; PS00070; ALDEHYDE_DEHYDR_CYS; UNKNOWN_1.
Yersinia pestis.
Bacteria; Proteobacteria;
                                    FHUB OR YP03390.
                                                    Ferrichrome transport system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Haselkorn R., Kyrpides N., Overbeek R.;
"The genome sequence of the facultative intracellular pathogen
Brucella melitensis.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Delveconio V.G., Kapatral V., Redkar R.J., Patra G., M. Ivanova N., Anderson I., Bhattacharyya A., Lykidis A., Jablonski L., Larsen N., D'Souza M., Bernal A., Mazur M. Selkov E., Elzer P.H., Hagius S., O'Callaghan D., Lete: Haselkorn B. Kurnidon N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN=16M / ATCC
MEDLINE=20020109;
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01-JUN-2002 (TrEMBLrel.
Hypothetical protein BME
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NCBI_TaxID=29459;
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Brucella melitensis
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                                                                                                                                            PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                        Conservative
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PubMed=11756688;
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Pred. No.
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   subdivision; Enterobacteriaceae;
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Result
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Maximum Match
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Maximum DB seq length: 2000000000
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Perfect score:
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US-10-176-918-380
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US-10-176-273-132
US-10-066-494-132
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RESULT 1
US-09-766-396-26
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Patent No. US20020013456A1
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Gregor
                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 105 amino ac
                                               REFERENCE/DOCKET NUMBER: 22908-0002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-7041
                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/857,389
FILING DATE: <UNKNOWN>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS: THE SCRIPPS RESEARCH INSTITUTE
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. US20020013456A1th Torrey Pines Road, TPC-8
                                                                                                                                                                                                                   ZIP: 92037
COMPUTER READABLE FORM:
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Siggins, George R.
Henriksen, Steven J.
TITLE OF INVENTION: CORTISTATIN:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/766,396 FILING DATE: 18-Jan-2001 CLASSIFICATION: <Unknown>
                                       TELEFAX: (415)
                                                                              NAME: Schmonsees, William REGISTRATION NUMBER: 31,796
                                                                                                                                                                                                                                      COUNTRY: US
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STATE: Califori
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US-10-066-211-132

US-10-142-419-380

US-10-123-262-380

US-10-142-423-380

US-10-141-755-380

US-10-141-755-380
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US-10-123-236
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US-10-140-921
US-10-140-928
US-10-123-292
US-10-123-903
US-10-124-822
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US-10-123-908
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RESULT 2
US-10-062-375-26
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Patent No. US20020133000A1
GENERAL INFORMATION:
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                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                               STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
ERAGMENT TYPE: internal
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TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: internal
SEQUENCE DESCRIPTION: SEQ I
                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/062,375
FILING DATE: 30-Jan-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 324-7041
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 MPLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/857,389 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: La Jolla
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
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                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                              NAME: Schmonsees, William REGISTRATION NUMBER: 31,796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 92037
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STREET: 10666 No. US20020133000A1th Torrey Pines Road, TPC-8
                                                                                                                                                              LENGTH: 105 amino acids
                                                                                                                                                                                                                     TELEFAX: (415)
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                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION DATA:
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                                                                                                                                                                                    CHARACTERISTICS:
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Siggins, George R.
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Pred. No. 5.4e-52;
Mismatches 0;
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US-10-066-500-132
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/066,500 CURRENT FILING DATE: 2002-02-01 PRIOR APPLICATION NUMBER: 10/002,796 PRIOR FILING DATE: 2001-11-15 PRIOR APPLICATION NUMBER: 60/056974 PRIOR APPLICATION NUMBER: 60/056974 PRIOR FILING DATE: 1997-08-26 PRIOR FILING DATE: 1997-08-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3130R1C7
                                                   PRIOR APPLICATION NUMBER: 60/066364
PRIOR FILING DATE: 1997-11-21
                                                                                       PRIOR FILING DATE: 1997-10-29
                                                                                                         PRIOR APPLICATION NUMBER: 60/063733
                                                                                                                                             PRIOR APPLICATION NUMBER: 60/063329
                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-10-24
                                                                                                                                                                                 PRIOR
                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/059263 PRIOR FILING DATE: 1997-09-18
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                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/PRIOR FILING DATE: 1997-10-17
                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/059588 PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60
PRIOR FILING DATE: 1997-09-1
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                                    APPLICATION NUMBER: 60/066840
                                                                                                                                                              APPLICATION NUMBER: 60/063082 FILING DATE: 1997-10-31
                   FILING DATE:
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Kevin P. Baker
David A. Botstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mary E. Gerritsen
Audrey Goddard
Paul J. Godowski
Austin L. Gurney
Ivar J. Kljavin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MPLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA
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P.Mickey Williams
William I. Wood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zemin Zang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Daniel Tumas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Timothy A. Stewart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Margaret Ann Roy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mary A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jennie P. Mather
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sherman Fong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Napoleone Ferrara
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Dan L. Eaton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hanspeter Gerber
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Pred. No. 5.4e-52;
; Mismatches 0;
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OR APPLICATION NUMBER: 60/079294
OR FILING DATE: 1998-03-25
OR APPLICATION NUMBER: 60/081049
OR FILING DATE: 1998-04-08
OR APPLICATION NUMBER: 60/095998
OR APPLICATION NUMBER: 60/095998
OR FILING DATE: 1998-08-10

APPLICATION NUMBER: FILING DATE: 1998-0

1998-02-09 1998-02-09 1997-12-16

APPLICATION FILING DATE:

NUMBER:

60/074086 60/074092

FILING DATE: 1998-09-10
APPLICATION NUMBER: 60/099811
FILING DATE: 1998-09-10
APPLICATION NUMBER: 60/099812
FILING DATE: 1998-09-10

APPLICATION NUMBER:

NUMBER: 60/099803 1998-09-10

APPLICATION NUMBER: 60/099601 FILING DATE: 1998-09-09

APPLICATION NUMBER: FILING DATE: 1998-0

1998-08-18

60/097000

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Query Match
Best Local Similarity
Matches 105; Conserv
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OR FILING DATE: 1998-09-10
OR APPLICATION NUMBER: PCT/US98/1903
OR FILING DATE: 1998-09-14
OR APPLICATION NUMBER: PCT/US98/19330
OR FILING DATE: 1998-09-15
OR APPLICATION NUMBER: PCT/US98/19437
OR FILING DATE: 1998-09-17
OR FILING DATE: 1998-09-17
OR APPLICATION NUMBER: PCT/US98/24855
OR APPLICATION NUMBER: PCT/US98/25108
OR APPLICATION NUMBER: PCT/US98/25108
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OR APPLICATION NUMBER: 09/870574
OR FILING DATE: 2001-05-30
OR APPLICATION NUMBER: 09/872035
OR FILING DATE: 2001-06-01
OR APPLICATION NUMBER: 09/886342
OR FILING DATE: 2001-06-19
OR APPLICATION NUMBER: PCT/US98/14552
OR FILING DATE: 1998-07-14
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DR FILING DATE: 1999-10-18
DR APPLICATION NUMBER: 09/403297
DR FILING DATE: 1999-10-18
DR APPLICATION NUMBER: 09/423741
DR FILING DATE: 1999-11-10
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DR APPLICATION NUMBER: 09/665350

R FILING DATE: 2000-09-18

DR APPLICATION NUMBER: 09/709238

DR FILING DATE: 2000-11-08

DR APPLICATION NUMBER: 09/767609

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DR FILING DATE: 2001-01-22
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R APPLICATION |
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APPLICATION NUMBER: PCT/US99/20111
FILING DATE: 1999-09-01
APPLICATION NUMBER: PCT/US99/20594
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FILING DATE: 2000-04-13
APPLICATION NUMBER: 09/664610
                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PCT/US98/25190 FILING DATE: 1998-11-25
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APPLICATION NUMBER: 09/522342
                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US99/12252
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FILING DATE: 1999-06-14
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100.
milarity 100.
Conservative
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NUMBER: PCT/US99/20594
: 1999-09-08
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                                                                                                               NUMBER: PCT/US99/21090: 1999-09-15
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DR FILING DATE: 1999-03-23

OR APPLICATION NUMBER: 60/139695

OR FILING DATE: 1999-06-16

OR FILING DATE: 1999-07-20

OR FILING DATE: 1999-07-26

OR APPLICATION NUMBER: 60/145698

OR FILING DATE: 1999-07-26

OR APPLICATION NUMBER: 60/149396

OR FILING DATE: 1999-08-17

OR PILING DATE: 1999-12-07

OR PILING DATE: 1999-12-07

APPLICATION NUMBER: 60/109304 FILING DATE: 1998-11-20 APPLICATION NUMBER: 60/125778

FILING DATE: 1998-10-APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-09 APPLICATION NUMBER: 60 FILING DATE: 1998-09-

1998-09-24

60/106032 60/101922 60/100858 9-17

DR APPLICATION NUMBER: 09/136828
DR FILLING DATE: 1998-08-19
DR APPLICATION NUMBER: 09/158342
DR FILLING DATE: 1998-09-21
DR APPLICATION NUMBER: 09/180997
DR FILLING DATE: 1998-09-10
DR FILLING DATE: 1998-09-10
DR APPLICATION NUMBER: 09/202088
DR FILLING DATE: 1998-12-08
DR APPLICATION NUMBER: 09/254311
DR FILLING DATE: 1999-03-03

R FILING DATE: 1999-04
R APPLICATION NUMBER:
R FILING DATE: 1999-06
R FILING DATE: 1999-06

1999-06-14

APPLICATION NUMBER: 09 FILING DATE: 1999-04-1

09/284663

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NUMBER: 09/254465: 1999-03-05

APPLICATION NUMBER:

09/254460

FILING DATE: 1999-03-09

DR APPLICATION NUMBER: 08/933821
DR FILING DATE: 1997 -09-19
R APPLICATION NUMBER: 08/960507
DR FILING DATE: 1997-10-29
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DR FILING DATE: 1998-07-14
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DR FILING DATE: 1998-08-19
DR FILING DATE: 1998-08-19
DR FILING DATE: 1998-08-19

APPLICATION NUMBER: 08/9 FILING DATE: 1997-08-26

08/918874

APPLICATION NUMBER: 09/ FILING DATE: 1998-08-19

09/136804

Score 554; DB 9; Pred. No. 8.4e-52; ; Mismatches 0; Length 155; Indels 0; Gaps

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RESULT 4
US-10-028-072-380
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CURRENT APPLICATION NUMBER: US/10/028,072
CURRENT FILING DATE: 2001-12-19
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OR FILING DATE: 1997-09-17
OR APPLICATION NUMBER: 60/059115
OR FILING DATE: 1997-09-17
OR APPLICATION NUMBER: 60/059117
OR APPLICATION NUMBER: 60/059117
OR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/059263
FILING DATE: 1997-09-18
APPLICATION NUMBER: 60/059352
FILING DATE: 1997-09-19
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APPLICATION NUMBER: 60/059588
FILING DATE: 1997-09-19
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                                  APPLICATION NUMBER: 60/
FILING DATE: 1997-10-24
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FILING DATE: 1997-09-24
APPLICATION NUMBER: 60/062250
FILING DATE: 1997-10-17
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APPLICATION NUMBER: 60/063082 FILING DATE: 1997-10-31
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FILING DATE: 1997-10-24
                                                                                                       APPLICATION NUMBER: 60/
FILING DATE: 1997-10-24
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                                                                                                                                                                              APPLICATION NUMBER: 60 FILING DATE: 1997-10-1
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FILING DATE: 1997-09-17
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Gurney, Austin L.
Sherwood, Steven
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Godowski, Paul
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Gerritsen, Mary E.
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DeForge, Laura
Desnoyers, Luc
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APPLICATION NUMBER: 60/064809
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APPLICATION NUMBER: 60/084600 FILING DATE: 1998-05-07

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Prior Application removed - See File
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> GENERAL INFORMATION: Sequence 380, A Publication No.

Application US/10123904 o. US20030022328A1

Baker, Kevin

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APPLICANT: I APPLICANT: APPLICANT: APPLICANT:

DeForge, Laura Beresini, Maureen

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Gerritsen,Mary E.

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APPLICANT

Stewart, Timothy A Sherwood, Steven Goddard, Audrey Godowski, Paul J. Gao, Wei-Qiang Desnoyers, Luc Filvaroff, Ellen

Smith, Victoria Gurney,Austin L.

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US-10-140-470-380
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US-10-123-904-380
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LENGTH: 155
TYPE: PRT
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LENGTH: 155
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                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/140,470 CURRENT FILING DATE: 2002-05-06
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NUMBER OF SEQ ID NOS: 550
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CURRENT FILING DATE: 2002-04-16
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Watanabe, Colin K
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Pred. No. 8.4e-52;
Pred. No. 8.4e-52;
                                                                             Score 554; DB 9;
Pred. No. 8.4e-52;
                                                                Mismatches
                                                                                              Length 155;
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US-10-176-918-380; Sequence 380, App. Publication No.
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Best Local :
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
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CURRENT APPLICATION NUMBER: US/10/175,746
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TYPE: PRT
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                                                                              Godowski, Paul J. Gurney, Austin L.
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Watanabe, Colin K
Wood, William
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Godowski, Paul J.
Gurney, Austin L.
                              Stewart, Timothy A
                                              Smith, Victoria
                                                               Sherwood, Steven
                                                                                                                Goddard, Audrey
                                                                                                                               Gerritsen, Mary E
                                                                                                                                              Gao, Wei-Qiang
                                                                                                                                                               Filvaroff, Ellen
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Watanabe,Colin K
                Tumas,Daniel
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Filvaroff, Ellen
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b. US20030027270A1
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Pred. No. 8.4e-52;
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111 SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 155

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LENGTH: 155
TYPE: PRT
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Best Local
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Best Local :
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SEQ ID NO 380
LENGTH: 155
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CURRENT APPLICATION NUMBER
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NUMBER OF SEQ ID NOS: 550
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CURRENT FILING DATE: 2002-06-20
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CURRENT FILING DATE: 2002-06-20
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Gurney, Austin L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DeForge, Laura
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o. US20030027276A1
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                                                                                                             Score 554; DB 9;
Pred. No. 8.4e-52;
Mismatches 0;
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RESULT 11
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PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059115
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CURRENT APPLICATION NUMBER: US/10/002,796
CURRENT FILING DATE: 2001-11-15
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APPLICATION NUMBER: 60/095998
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P.Mickey Williams
William I. Wood
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Timothy A. Stewart
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 0;
Score 554; DB 9;
Pred. No. 8.4e-52;
; Mismatches 0;
                         Length 155;
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RESULT 12
US-10-066-273-132
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CURRENT FILING DATE: 2002-02-01
PRIOR APPLICATION NUMBER: 10/02,796
PRIOR FILING DATE: 2001-11-15
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
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PRIOR FILING DATE: 1997-09-17
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             OR APPLICATION NUMBER: 60/079294

RE FILLING DATE: 1998-03-25

DR APPLICATION NUMBER: 60/081049

DR FILLING DATE: 1998-04-08

DR APPLICATION NUMBER: 60/095998

DR APPLICATION NUMBER: 60/095998

DR FILLING DATE: 1998-08-10
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APPLICATION NUMBER: 60/074092
FILING DATE: 1998-02-09
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FILING DATE: 1997-10-27
APPLICATION NUMBER: 60/063733
FILING DATE: 1997-10-29
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P.Mickey Williams
William I. Wood
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Timothy A. Stewart
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Hanspeter Gerber
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David A. Botstein
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OR APPLICATION NUMBER: 09/333075
IR FILING DATE: 1999-06-14
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OR APPLICATION NUMBER: 09/380137
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OR FILING DATE: 1999-08-25 OR FILING DATE: 1999-07-20

R APPLICATION NUMBER: 60/145698

OR FILING DATE: 1999-07-26

OR APPLICATION NUMBER: 60/149396

OR FILING DATE: 1999-08-17

OR APPLICATION NUMBER: 60/169495 R FILING DATE:
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APPLICATION NUMBER: 09/284663
FILING DATE: 1999-04-15
APPLICATION NUMBER: 09/332928 APPLICATION NUMBER: 09/254311 FILING DATE: 1999-03-03 APPLICATION NUMBER: 09/254460 FILING DATE: 1999-03-09 APPLICATION NUMBER: 09/254465 FILING DATE: 1998-08-19 APPLICATION NUMBER: 09/ FILING DATE: 1998-08-19 APPLICATION NUMBER: 08/960507 FILING DATE: 1997-10-29 APPLICATION NUMBER: 09/114844 FILING DATE: 1998-07-14 APPLICATION NUMBER: 09/136801 FILING DATE: 1999-12-07
APPLICATION NUMBER: 08/918874
FILING DATE: 1997-08-26
APPLICATION NUMBER: 08/93821
FILING DATE: 1997-09-19 APPLICATION NUMBER: 60/101922 FILING DATE: 1998-09-24 APPLICATION NUMBER: 60/106032 APPLICATION NUMBER: 09/136828 FILING DATE: 1998-08-19 APPLICATION NUMBER: 09/158342 FILING DATE: 1998-09-21 APPLICATION NUMBER: FILING DATE: 1999-0 APPLICATION NUMBER: 60/100858 APPLICATION NUMBER: 09/332929 APPLICATION NUMBER: 09/202088 APPLICATION NUMBER: 09/180997 APPLICATION NUMBER: 60/145070 APPLICATION NUMBER: 60/125778 APPLICATION NUMBER: 60/109304 FILING DATE: 1998-11-20 FILING DATE: 1998-10-FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 1998-09-09 APPLICATION NUMBER: 60/ FILING DATE: 1999-06-14 FILING DATE: FILING DATE: FILING DATE: 1998-09-10 APPLICATION NUMBER: 60/099601 FILING DATE: 1998-09-10 FILING DATE: NUMBER: 09/380139 : 1999-08-25 NUMBER: 09/403296 1999-06-1 1998-12-08 1999-06-15 1999-03-23 1998-09-1998-09-10 1999-10-1 09/380138 09/136804 08/960507 60/099812 60/139695 60/099811 60/099803

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FILING DATE: 1998-12-01
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APPLICATION NUMBER: 09/665350
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APPLICATION NUMBER: 09/423741
FILING DATE: 1999-11-10
APPLICATION NUMBER: 09/423844
FILING DATE: 1999-11-12
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FILING DATE: 1999-06-02
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Pred. No. 8.4e-52;
Mismatches 0;
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Sequence 132, Application US/10066494

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PRIOR FILING DATE: 2001-11-15
PRIOR FILING DATE: 2007-11-15
PRIOR PAPELICATION NUMBER: 60/05974
PRIOR PAPELICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR PELICATION NUMBER: 60/059263
PRIOR PELICATION NUMBER: 60/059263
PRIOR APPLICATION NUMBER: 60/059588
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TITLE OF INVENTION: SCRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3130R1C9
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OR APPLICATION NUMBER: 60/066840
OR FILING DATE: 1997-11-25
IOR APPLICATION NUMBER: 60/069694
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APPLICATION NUMBER: 60/062816
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FILING DATE: 1998-04-08
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f: Navid A. Botstein

havid A. Tovers
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P.Mickey Williams
William I. Wood
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Margaret Ann Roy
Timothy A. Stewart
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Hanspeter Gerber
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DR APPLICATION NUMBER: 09/284663
DR FILING DATE: 1999-04-15
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DR FILING DATE: 1999-06-14
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APPLICATION NUMBER: 09/333075
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FILING DATE: 1998-12-08
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APPLICATION NUMBER: 09/158342
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FILING DATE: 1999-12-07
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APPLICATION NUMBER: 60/
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FILING DATE: 1999-06-14
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FILING DATE: 1999-03-23
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Best Local :
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R APPLICATION NUMBER: PCT/US99/05028

OR FILING DATE: 1999-03-08

OR APPLICATION NUMBER: PCT/US99/12252

OR FILING DATE: 1999-06-02

OR FILING DATE: 1999-09-01

OR FILING DATE: 1999-09-01
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APPLICATION NUMBER: 09/665350
TING DATE: 2000-09-18
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FILING DATE: 1998-11-20
APPLICATION NUMBER: PCT/US98/25108
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APPLICATION NUMBER: PCT/US98/25190
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APPLICATION NUMBER: PCT/US98/19093
FILING DATE: 1998-09-14
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APPLICATION NUMBER: 09/886342
FILING DATE: 2001-06-19
APPLICATION NUMBER: PCT/US98/14552
FILING DATE: 1998-07-14
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FILING DATE: 2000-11-08
APPLICATION NUMBER: 09/767609
FILING DATE: 2001-01-22
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APPLICATION NUMBER: 09/870574
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61 SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
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                                                                                                                                                                                                                          NUMBER: PCT/US99/21547
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Pred. No. 8.4e-52;
Mismatches 0;
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RESULT 14 US-10-137-865-380

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Sequence 380, Applicatic Publication No. US200300 GENERAL INFORMATION: APPLICANT: Baker, Kevin Application US/10137865 o. US20030032155A1

APPLICANT: APPLICANT

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APPLICANT:

Beresini, Maureen DeForge, Laura

Desnoyers, Luc Filvaroff, Ellen

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US-10-137-865-380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-140-474-380
Prior Application removed -
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 380
LENGTH: 155
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 554; DB 9; Best Local Similarity 100.0%; Pred. No. 8.4e-52; Matches 105; Conservative 0; Mismatches 0;
                                                                     APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: 93330R1C162
CURRENT APPLICATION NUMBER: US/10/140,474
CURRENT FILING DATE: 2002-05-06
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C154
CURRENT APPLICATION NUMBER: US/10/137,865
CURRENT FILING DATE: 2002-05-03
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Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
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                                                                                                                                                                                Watanabe, Colin K
Wood, William
                                                                                                                                                                                                                    Stewart, Timothy A. Tumas, Daniel
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Godowski, Paul J.
                                                                                                                                                                                                                                                                                                                                                                  Gao, Wei-Qiang
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Filvaroff, Ellen
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                                                                                                                                                                                                                                                           Smith, Victoria
                                                                                                                                                                                                                                                                          Sherwood, Steven
                                                                                                                                                                                                                                                                                             Gurney, Austin L.
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                                                                                                                                                                                                                                                                                                                                                                                                                       DeForge, Laura
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Gerritsen,Mary E.
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Best Local :
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111 SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 155
                                                                                                                            105;
               61 SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                          51 MPLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA 110
                                                                             1 MPLSPGILLILLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA 60
                                                                                                                         h 100.0%;
Similarity 100.0%;
05; Conservative (
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Pred. No. 8.4e-52;
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Search completed: June 23, 2003, 16:06:11 Job time: 49 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Sequence:
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2003 Compus
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/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
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US-08-628-929-7

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US-09-218-950-27
                                          PCT-US92-01785-27
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US-09-039-116-1
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1,	Sequence 1, Appli	Sequence 1, Appli	Sequence 83, Appl	Sequence 1, Appli	1,	Sequence 83, Appl	Sequence 6, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	 Sequence 1, Appli 	Sequence 13, Appl	Sequence 1, Appli	Sequence 4, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli

ALIGNMENTS

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; MOLECULE TYPE: US-09-001-472-2
  Query Match
Best Local Similarity
Matches 105; Conserv
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APPLICATION NUMBER: US/UY/UL.
FILING DATE: Herewith
CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/033,980
APPLICATION NUMBER: US 60/033,980
APPLICATION NUMBER: US 60/033,980
                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                     NAME: STEFFE, ERIC K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/033,980
FILING DATE: 31-DEC-1996
PRIOR APPLICATION TOWNS:
APPLICATION NUMBER: US 60/037,386
FILING DATE: 07-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: STEFFE, ERIC K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 20005-3934

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
STREET: 1100 NEW YORK AVENUE, SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: OLSEN, HENRIK S.
APPLICANT: RUBEN, STEVEN M.
TITLE OF INVENTION: CORTISTATIN POLYPEPTIDES
                                                                                                                                                                                                                        TELEFAX:
                                                                                                                        TOPOLOGY:
                                                                                                                                                            LENGTH:
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STATE: DC
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                                                                                                                                     amino acid
  Conservative
                                                                                                                                                                                                                        (202)
                                                                                                                        linear
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371-2540
                  100.0%; Score 554; DB 4; 100.0%; Pred. No. 4.5e-62;
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                                                                                                                                                    Patent No.
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                                                                                                                       Sequence 3, Application of the 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (610) 554-6312
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 112 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC pc-po5/mS-Dos
OPERATING SYSTEM: PC-DO5/mS-Dos
                                                                          APPLICANT:
NUMBER OF SEQUENCES: 1
                                                  TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 554-2937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Sutcliffe, APPLICANT: de Lecea, 1
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                                                                               RUBEN,
                                                                                                    OLSEN,
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STEVEN M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           34,163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 295; DB 3;
Pred. No. 1.5e-29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         519.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 112;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (202) 371-25
INFORMATION FOR SEQ ID NO:
                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 31-DEC-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/001,472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                       CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 07-FEB-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                            STREET: LVC.
                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                  SOFTWARE:
                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 36 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 60 FILING DATE: 31-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             No. 6074872
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STEFFE, ERIC K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/08648322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SSTAFEGGTPELSKRQERPPLQQPPHRDKKPCKNFFWKTFSSCK 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PSALSLLLLLLLSGIAASALPLESGPTGQDS--VQDATGGRRTGLLTFLAWWHEWASQDS 68
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de Lecea, L
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                                               PatentIn Release #1.0, Version #1.25
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SYSTEM: PC-DOS/MS-DOS
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COMPOSITIONS
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                                                                                                                                                                                                                                                                                                                                                                                AND METHODS
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Gaps

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US-08-648-322-6
   Query Match
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                               MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 51
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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APPLICANT: de Lecea, l
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CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                              TOPOLOGY:
                                                                                                                                                                         TELEPHONE:
                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                    SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
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                                                                                                            ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                   92037
                                                                                            amino acid
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                                                                                                            85 amino acids
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                                                                                                                                                       (619) 554-6312
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                                                                                                                                                                           (619) 554-2937
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C-terminal
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   48.2%;
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   Score 267;
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Length 85;
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Sequence 4, Application US/09001472
Patent No. 6232100
GENERAL INFORMATION:
APPLICANT: OLSEN, HENRIK S.

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RESULT 7
US-09-001-472-4
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US-08-648-322-10
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                                                                                                                                                                                                     Query Match
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Matches 5
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                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                 MOLECULE TYPE: FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 84 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: PC-D
SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Fitting, Thomas REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
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                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                          LENGTH:
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                                                                                     79 GAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                                                                                                             19 AALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQASAGPLIGEEAREVARRQE 78
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 50; Conserv
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: California
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                                                         RPPPQQPPHLDKKPCKNFFWKTFSSCK
                                                                                                                  SALPLESGPTGQDS--VQEATEGR-SGLLTFLAWWHEWASQASSSTPVGGGTPGLSKSQE 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/08648322
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SYSTEM: PC-DOS/MS-DOS
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C-terminal
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                                                                                                                                                                                      Score 257.5; DB Pred. No. 5e-25;
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                                                         84
                                                                                                                                                                                                    DB 3;
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                                                                                                                                                                          Indels
                                                                                                                                                                                                     Length
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                                                                                                                                                                        Gaps
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US-08-648-322-5

MOLECULE TYPE: protein

TOPOLOGY: TYPE:

amino acid

linear

TELEFAX: (619) 554-63 INFORMATION FOR SEQ ID NO:

TELEPHONE:

TELECOMMUNICATION INFORMATION:

(619)

554-2937

554-6312

<u>ت</u>

REFERENCE/DOCKET NUMBER:

ATTORNEY/AGENT INFORMATION:

CLASSIFICATION:

FILING DATE:

NAME: Fitting, Thom: REGISTRATION NUMBER:

Thomas

34,163

519.0

SEQUENCE CHARACTERISTICS: LENGTH: 109 amino acid

109 amino acids

Matches Query Match

58;

Conservative

11;

Mismatches

29;

6;

Gaps

61

12

2 PLSPGLLLLLLSGATATAAALPLEGGPTGRDSEHMQEAAGIRKSSLLTELAWWFEWTSQAS

PSAFGLLLL---WGVAASALPLESGPTGQDS--VQEATEGR-SGLLTFLAWWHEWASQAS 65

Local

Similarity

49.1%;

Score 272; Pred. No. 1.

DB 3; .1e-26;

Length 109; Indels

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                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                              yeurce 11, Application US/08648322 itent No. 6074872
              CURRENT APPLICATION DATA:
                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. 6074872th Torrey Plnes Road, TPC-8
                                                                                                                                                                                                                                                                     APPLICANT: Sutcliffe, Gregor J. APPLICANT: de Lecea, Luis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: RUBEN, ST
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 60/033,980 FILING DATE: 31-DEC-1996 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 60/037,386 FILING DATE: 07-FEB-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 60/033,980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/001,472
FILING DATE: Herewith
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                                                                                                                                          CITY: La Jolla
STATE: California
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 APPLICATION NUMBER:
                               SOFTWARE:
                                                                                                                            COUNTRY:
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                                                                                                              92037
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)GY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WASHINGTON
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                               PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                                                                                      CORTISTATIN: NEUROPEPTIDES, COMPOSITIONS AND METHODS
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US/08/648,322
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Pred. No. 2e-13;
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                                          Query Match
              Matches
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                                                                                                                                        TYPE:
                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
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              Conservative
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; MOLECULE TYPE: protein ; FRAGMENT TYPE: C-terminal US-08-648-322-11
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Best Local Similarity 03...
20; Conservative
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Patent No. 6074872
                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                       TELEFAX: (619) 554-63 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (619) 554-6312 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 51
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 554-2937
                                                                                  MOLECULE TYPE: protein FRAGMENT TYPE: C-terminal
                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 554-2937
                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/0
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                         NAME: Fitting, Thomas REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    77 QEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
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                                                                                                                                    amino acid
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)GY: linear
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                                                                                                                                                       29 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: THE SCRIPPS RESEARCH INSTITUTE
10666 No. 6074872th Torrey Pines Road, TPC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29 amino acids
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de Lecea, l
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                                                                                                                     linear
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554-6312
NO: 7:
               20.9%;
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                                                                                                                                                                                                                                                                                                                                                                                          Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                      34,163
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               Score 116; DB 3; Length 29; Pred. No. 6.6e-08;
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Pred. No. 2.8e-08;
Mismatches
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Indels
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Gaps
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; MOLECULE TYPE:
; FRAGMENT TYPE:
'US-08-648-322-3
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US-08-648-322-3
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                                                                                                                                                        Sequence 8, Application US/08648322 Patent No. 6074872
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (619) 554-2937
TELEFAX: (619) 554-6312
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
                                   TITLE OF INVENTION: CO
TITLE OF INVENTION: CO
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: 51
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: THE SCRIPPS RESEARCH INSTITUTE
STREET: 10666 No. 6074872th Torrey Pines Road, TPC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
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NAME: Fitting, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH:
                         DDRESSEE:
                                                                                                                                                                                                                                                            63 EPEDLPQAAEQDEMRLELQRSANSNPAMAPRERKAGCKNFFWKTFTSC 110
                                                                                                                                                                                                                                                                                                                                                             17 ATAALPLE----GGPTGRDS-----EHMQE--AAGIRKSSLLT-FLAWWFEWTSQASAGPL
                                                                                                                                                                                                                                                                                           66 IGEEAREVARROE------GAPPQQSARRDRMPCRNFFWKTFSSC 104
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                                                                                                                                                                                                                                                                                                                                                                                                      34;
                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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THE SCRIPPS RESEARCH INSTITUTE 10666 No. 6074872th Torrey Pines Road, TPC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       110 amino acids
                                                                                    Sutcliffe, Gregor J.
de Lecea, Luis
VENTION: CORTISTATIN:
                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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de Lecea, Luis
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C-terminal
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                                                                     COMPOSITIONS AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                 17.1%; Score 95;
31.5%; Pred. No.
tive 13; Mismatc
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                                                                                       NEUROPEPTIDES,
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                                                                                                                                                                                                                                                                                                                                                                                                                   0.00017;
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   SEQUENCE CHARACTERISTICS:
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RESULT 12
US-08-648-322-23
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Best Local Similarity
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APPLICANT: Sutcli
APPLICANT: de Lec
   INFORMATION FOR SEQ
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                            NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: 51
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 554-2937
                                                                                                                                                                                                                                                                                     STREET: LOUG
STREET: LA Jolla
CTATE: California
                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
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FRAGMENT TYPE:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                TELEFAX:
                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
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California
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ENTION: CORTISTATIN:
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C-terminal
554-6312
ID NO: 2
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                            Sequence 8, Application US/09423890 Patent No. 6312934
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 APPLICANT:
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                                                                                                                                                                                                                                  TOPOLOGY: 11:
MOLECULE TYPE:
FRAGMENT TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
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FRAGMENT TYPE:
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                                                                                                                                                                 Local Similarity les 12; Conserv
                                                                                                                                                                                                                                                                         TYPE: amino acid
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               INFORMATION:
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XX: linear
                                                                                                            PCKNFFYKTFSSCK 14
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                                                                                                                                                                 Conservative
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de Lecea, Luis
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C-terminal
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C-terminal
PHARMACEUTICAL CORPORATION
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85.7%;
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Pred. No.
                                                                                                                                                                               Score 76;
Pred. No.
                                                                                                                                                               Mismatches
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Search completed: June 23, Job time: 27 secs
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US-08-628-829-4
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                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn
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NUMBER OF SEQ ID NOS: 25
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EARLIER APPLICATION NUMBER: 08/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 1996-04-05
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PRIOR FILING DATE: 1998-09-04
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PRIOR FILING DATE: 1998-03-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EARLIER FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EARLIER APPLICATION NUMBER: 08/323,460
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                                                                                                                                                                                                                                                                                                                                   LENGTH: 1593
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SARLIER APPLICATION NUMBER: 08/410,602
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      !ITLE OF INVENTION: Method And Product For Regulating Cell Responsiveness To Exter
!ILE REFERENCE: CPI-004DVCP3
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                                                                          298 ERRNRRGPVVVKPIPIKGDGSEVNNLAAEPQGEGQAGSAAPAPKGRRSPSP 348
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                                                                                                               58
                                                                                                                                                                                    13 SGATATAAL--PLEGGPTGRDSEHMQEAAGIRKS-----SLLTFLAWWFEWT 57
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                                                                                                                                                 SGARSPAGAEPPSAAAPSGREMENKETLKGLHKMEDRPEERMIREKLKATCMPAWKHEWL 297
                                                                                                                                                                                                                                                                                                                    Mus musculus
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                                                                                                             -GEEAREVARRQEG-----APPQQSARRDRMP 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -GEEAREVARRQEG-----APPQQSARRDRMP
                                                                                                                                                                                                                                           Score 71.5; D
Pred. No. 5.7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 71.5; D
Pred. No. 5.2;
                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                          DB 4;
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                                                                                                                                                                                                                                                          Length 1593;
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                                                                                                                                                                                                                      Indels 31;
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Result
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 otal number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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554
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Listing first 45 summaries
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Copyright (c) 1993 - 2003 Compugen Ltd.
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85
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                                                                                                                                                                                                                                                                                         SUMMARIES
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                                                                                                                                                                                                                         Description
  Amino a
Novel h
hCS-85
Mutant
hCS-62
                                                                                                                                                                Human cortistatin hCS-105 peptide.
                                                                                                     Human
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17.9	17.9		80	80	18.2	•	18.8	18.8	18.8	18.8	20.0	20.9	•	21.5	•	•	•	28.0	•	•	•	28.9	•	•	•	•		•	•	80	9	9.	53.2	53.2
16	16	16	17	17	17	17	17	17	17	17	29	29	29	29	29	23	51	28	28	28	29	29	29	29	29	33	84	85	84	85	109	109	112	112
23	21	19	23	21	19	19	23	22	22	19	22	23	19	23	19	19	23	19	19	19	19	19	19	19	19	19	23	23	19	19	23	19	23	19
ABB73263	AAB17373	AAW44021	ABB73272	AAB17382	AAW44024	AAW44030	ABB73260	AAB91000	AAB90999	AAW44016	AAB91946	AAE21874	AAW42036	AAE21878	AAW42040	AAW44054	AAE21924	AAW44051	AAW44050	AAW44049	AAW44048	AAW64434	AAW44046 .	AAW44042	AAW44047	AAW44053	AAE21877	AAE21873	AAW4 2039	AAW4 2035	AAE21872	AAW42034	8	AAW42033
atostatin/co	agon i	Mutated hCS peptid	tatin/co	agoni	hcs	Mutated hCS peptid	tatin	Somatostatin relat	Somatostatin relat	hCS-17 peptide. S	Cortistatin peptid	Rat procortistatin	A fragment of rat	Mouse procortistat		Mutant hCS peptide	Η		hcs-29	-29	S	tic cort	nt hCS-29 pep	peptide	hcs-	t hCS peptid	Mouse procortistat	Rat procortistatin	. Mouse procortistat	Rat procortistatin		Mouse preprocortis	t preprocortis	Rat preprocortista

ALIGNMENTS

RESULT 1

AAW64433 standard; Protein; 105 AA.

Cortistatin; human; diagnosis; Human cortistatin protein.

treatment; sleep disorder

16-OCT-1998 AAW64433;

(first entry)

07-FEB-1997; 31-DEC-1996; Key Peptide Olsen HS, Protein 23-DEC-1997; 09-JUL-1998. Homo sapiens (HUMA-) HUMAN GENOME SCI INC WO9829438-A2 Ruben SM; 97US-0037386. 96US-0033980. 97WO-US23784 /label= signal 20..105 /label= cortistatin Location/Qualifiers

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Best Local S
Matches 105
                                                                                                                                                                                                          15-OCT-1996;
07-JUN-1996;
19-SEP-1996;
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N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence represents a novel human cortistatin which can be in diagnostic methods for detecting variations in cortistatin generation. The cortistatin polypeptide can be used for treating an individual in need of an increased level of cortistatin e.g. diagnosis and treatment of a sleep disorder.
                                                                                                                                                                                                                                                                                                                                          Corticostatin; somatostatin; hCS-105 peptide; hormone-producing tumour; gastric ulcer; dementia; growth disorder; hormone secretion regulation; digestive system regulation; neural inhibitor; therapy.
 6
         This sequence represents the peptide hCS-105. It is a peptide of the invention, and has corticostatin or somatostatin activity. Antibodies recognising hCS-17 can be used to screen for a compound that modulates, i.e. an agonist or antagonist, the binding of hCS-17 to its receptor, a
                                                                  Claim
                                                                                      growth
                                                                                                Peptide having corticostatin or anticancer and antiulcer agent,
                                                                                                                                  N-PSDB;
                                                                                                                                                                   Fukusumi
                                                                                                                                                                                       (TAKE ) TAKEDA CHEM IND
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                                                                                                                                 1998-042177/04
DB; AAV02054, AJ
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                                                                                      abnormalities
                                                                                                                                                                                                                                                                                                                                                                                        peptide
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                                                                                                                                                                 s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MPLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fig 1; 81pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MPLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA
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                                                                                                                                                                 Hinuma
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                                                                116; 174pp;
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96JP-0146052.
96JP-0247710.
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                                                                                                                                  AAV02055
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                                                                 Japanese
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Pred.
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                                                                                                 somatostatin activity - use and for control of dementia
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.2e-54;
s 0;
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RESULT 3
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                                          This sequence encodes the novel human preprocortistatin protein, which is processed to produce the mature cortistatin proteins referred as human cortistatin-29, and cortistatin-17. The purified cortistatin, and its agonists, are used to induce sleep while its receptor antagonists (particularly antibodies) is used to inhibit sleep. Although cortistatin is structurally similar to somatostatin, it is able to depress neuronal electrical activity, induce low frequency waves in the cerebral cortex, and the cortistation is structurally similar to somatostatin, it is able to depress neuronal electrical activity, induce low frequency waves in the cerebral cortex,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hormone-producing tumours (e.g. tumours producing gastrin or insulin), gastric ulcers and dementia, regulate sleep and control growth disorders (e.g. acromegaly, gigantism and dwarfism). They can also be used to regulate hormone secretion and the digestive system (e.g. to treat diabetes), and as tumour multiplication or neural inhibitors.
                          electrical activity, induce low frequency waves in the cerebra antagonise acetylcholine and therefor enhance slow-wave sleep.
                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                                                                              mutation(s)
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                                                                                                                                                                                                                                                                                                           New cortistatin mutation(s) and
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st; antibody; inhibition; sleep; somatostatin; acetylcholine;
nal electrical activity; cerebral cortex; antagonist; primer; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preprocortistatin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hybridisation;
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                                                                                                                                                                                                                                                         112; 128pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96US-0648322
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                                                                                                                                                                                                                                                                                                              peptide(s)
screen for
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                                                                                                                                                                                                                                                           English.
                                                                                                                                                                                                                                                                                                           drugs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Siggins
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                                                                                                                                                                                                                                                                                                                                      used
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Pred. No. 8.2
); Mismatches
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.2e-54;
s 0;
                                                                                                                                                                                                                                                                                                                                      sleep,
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RESULT 4
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Best Local :
              The invention relates to a substantially isolated and purified mammalian cortistatin. The cortistatin nucleic acids, proteins, polypeptides and antibodies are useful in screening and diagnostic methods, and in therapeutic methods related to modulation of sleep and sleeping disorders. Cortistatin proteins may be used as immunogen to produce antibodies immunoreactive with cortistatin, in in vitro ligand binding specificities, to characterise candidate pharmaceutical compounds useful for modulating cortistatin functions. The present sequence is human effecting cortistatin functions. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and hybridisation/amplification assays to detect or quantify cortistatin (including that administered therapeutically) or its nucleic acid. Oligonucleotides, e.g. antisense molecules, are used in vivo to alter cortistatin gene expression. Detection of a mutation in the cortistatin gene may provide diagnosis of sleep-related or neuronal depression-related disorders or diseases of the brain.
                                                                                                                                                                                                                                                                                                          Claim 2; Fig 3b; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                        New mammalian cortistatin useful in screening and diagnostic methods, and in therapeutic methods related in modulating sleep and sleeping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sutcliffe JG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-MAY-1997;
15-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-JAN-2001; 2001US-0766396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-JAN-2002
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Cleavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human preprocortistatin protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAE21885;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAE21885 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SCRI ) SCRIPPS RES INST.
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DB; AAD34533, AAD34577.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; cortistatin;
disorder; pha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lecea LD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-0857389
96US-0648322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers 75..76 87..88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tin; sedative; neuropeptide; screening; therapeutic;
pharmaceutical compound; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Henriksen
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Pred. No. 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Siggins GR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          105;
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RESULT 5
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Best Local (
                                                                                                            20-MAR-2000;
21-MAR-2000;
30-MAR-2000;
17-MAY-2000;
         Baker KP,
Gerritsen
Smith V,
                                                                                                                                                      24-FEB-
24-FEB-
01-MAR-
                                                                                                                                                                                                                                                  16-DEC
20-DEC
20-DEC
30-DEC
                                                  (GETH
                                                                                                                                                                                                18-FEB
                                                                                                                                                                                                                                                                                                                                                                                                                                                   breast; prostate;
cartilage; ear; pr
adipocyte; A-pepti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                           06-JAN
                                                                                                                                                                                                                                                                                             09-DEC-1999;
                                                                                                                                                                                                                                                                                                                                               01-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                   01-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                            WO200140466-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAU12361;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAU12361 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                      11-FEB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   secretory and transmembrane;
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                                                  GENENTECH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
         Beresini M,
n ME, Goddard
Stewart TA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   ear; proliferation; glucose;
A-peptide; factor VIIA; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  105
                                                                    2000WO-US00277
2000WO-US00376
2000WO-US003565
2000WO-US04341
2000WO-US04344
2000WO-US04414
2000WO-US04914
2000WO-US05601
2000WO-US05601
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2000WO-US05601
2000WO-US057377
2000WO-US05732
2000WO-US057332
2000WO-US057332
2000WO-US05733705
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                                                                                                                                                                                                                                                                                                                                                                     2000WO-US32678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polypeptide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                            99WO-US28565.
99US-0170262.
99WO-US30095.
99WO-US30911.
99WO-US30999.
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                                                                                                                                                                                                                                                      99WO-US31243
                                                                                                                                                                                                                                                                                                                            99WO-US28551
                                                                                                                                                                                                                                                                                                                                                                                                                                                   nd transmembrane; PRO; mammalian; cancer; lung; cervical; tumour necrosis factor-alpha; TNF-alpha; roliferation; glucose; free fatty acid; skeletal muscle; ide; factor VIIA; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
      Deforge L, Desnoyers L, Filva
A, Godowski PJ, Gurney AL, Sh
Matanabe CK, Wood WI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 554; DB 23;
Pred. No. 8.2e-54;
; Mismatches 0;
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        Filvaroff E,
AL, Sherwood S
ood WI, Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 105;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC AAU12172-AAU12446 represent novel human secretory and transmembrane CC PRO polypeptides. The PRO polypeptides are useful to detect other CPRO polypeptides, to link bloactive molecules to cells expressing CC PRO polypeptides, to modulate biological activities of cells expressing CC PRO polypeptides, and to detect the presence of mammalian lung, colon, CC polypeptides, and to detect the presence of mammalian lung, colon, CC polypeptide expression in a cell sample to that in a control sample. CC Some of the 275 sequences are also useful to stimulate the release of tumour necrosis factor-alpha (TNF-alpha) from human blood, the CC proliferation or differentiation of chondrocytes, the proliferation or gene expression in pericyte cells, the release of proteoglycans from cartilage, the proliferation of inner ear utricular supporting cells or of T-lymphocytes, the release of a cytokine from peripheral blood CC monocytes (PBMCs), or the proliferation of endothelial cells. Some of the PRO polypeptides in a proliferation of endothelial cells. Some of the PRO polypeptides may modulate glucose or free fatty acid uptake by skeletal muscle cells or by adipocytes; or inhibit binding of A-peptide CC molecules involved in binding interactions. The polymucleotides encoding PRO polypeptides can be used to generate probes, antisense RNA/DNA, CC transgenic or knock out animals and can be used in gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                              Human; secreted protein; transmembrane protein; PRO196; PRO444; PRO183; PRO185; PRO210; PRO215; PRO217; PRO242; PRO288; PRO365; PRO1361; PRO1308; PRO1187; PRO1479; PRO449; PRO353; PRO1318; PRO1600; PRO940; PRO533; PRO301; PRO187; PRO337; PRO1411; PRO4356; PRO265; PRO941; PRO1096; PRO6003; PRO6004; PRO350; PRO2630;
                                                                                                                                                                                                                                                                                                                                                                                                                              AAB31213;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB31213 standard; Protein; 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated, secretory and transmembrane PRO polypeptide used to detect other PRO polypeptides, link bioactive molecules to cells expressing PRO polypeptides, and detect the presence of mammalian tumours e.g. lung, breast, prostate, cervical
                                                                          Modified-site
                                                                                                                   Modified-site
                                                                                                                                                                           Ното
                                                                                                                                                                                                                PRO6309;
                                                                                                                                                                                                                                                                                                                                               Amino acid sequence of human polypeptide PRO350.
                                                                                                                                                                                                                                                                                                                                                                                          20-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 12; Fig 380; 813pp; English.
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                                                                                                                                                                           sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                  cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MPLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA 110
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                                                                          /note=
35..39
                                                                                                                 Location/Qualifiers 33..37
                  51..69
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                                                                                                                                                                                                                genetic disorder;
                                  "cAMP- and cGMP-dependent protein kinase
phosphorylation site"
                                                                                             "amidation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 554; DB 22;
Pred. No. 1.3e-53;
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                                                                                                                                                                                                            transgenic animal; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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22-FEB-2000;
01-MAR-2000;
02-MAR-2000;
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Ferrara N, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
Godowski PJ, Gurney AL, Kljavin IJ, Mather JP, Napier MA, Pan J;
Paoni NF, Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM;
                                                                                                                                                                                       Isolated nucleic acid molecule encoding a PRO polypeptide which is a transmembrane polypeptide is useful for gene therapy and identification of related polypeptides -
                                                                                                                                                                                                                             WPI; 2001-050091/06.
N-PSDB; AAC87056.
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07-DEC-1999;
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Wood WI, Zhang Z;
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30-MAR-2000;
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18-FEB-2000;
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                                                                                                                                                                                                                                                                                                             (GETH )
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17-AUG-1999;
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                                                                                                                                                                                                                                                                                                             GENENTECH INC
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2000WO-US04341.

2000WO-US04414.

2000WO-US04414.

2000WO-US05601.

2000WO-US05841.

2000WO-US07377.

2000WO-US07377.

2000WO-US0843.

2000WO-US0843.

2000WO-US0843.
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99WO-US21547.
99WO-US28313.
99WO-US28301.
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99US-0169495.
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99US-0149396
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99US-0145070
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75..81
/note=
90..96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "N-myristoylation site"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "N-myristoylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   site'
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Claim 12; Fig 70; 244pp; English.

The present sequence represents a human secreted and transmembrane polypeptide. The specification describes human polypeptides, designated RR0196, pR0444, PR01183, PR0185, PR0210, PR0215, PR0217, PR0242, PR0288, PR0365, PR01311, PR01308, PR01183, PR01272, PR01419, PR04999, PR07170, PR0248, PR0353, PR01318, PR01600, PR09940, PR0533, PR0301, PR0187, PR0248, PR0353, PR01411, PR04356, PR0246, PR0265, PR0265, PR0301, PR06003, PR06004, PR0350, PR02630 and PR06309. The biological activity of cells can be modulated with agents that bind to these polypeptides, resulting in the death of the cells. The polypuclectides encoding these polypeptides are useful in the recombinant production of the polypeptides, as a hybridisation probe to screen libraries to isolate polypeptides, as a hybridisation probe to screen libraries to isolate polypeptides, as a hybridisation probe to screen libraries to isolate homologous sequences, or to map the gene. They may also be used for analysing genetic disorders, and to produce transgenic animals which are useful for the development and screening of therapeutically useful reagents. The polynuclectides can also be used in gene therapy e.g. to defective

Sequence 155

Q

100.0%; Score 554; B 22; Length

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189;

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RESULT 7
ABG22878
ID ABG2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to isolated polynucleotide (I) and CC polymerase (II) sequences. (I) is useful as hybridisation probes, CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome CC and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags CC for identifying expressed genes. (I) is useful in gene therapy techniques CC to restore normal activity of (II) or to treat disease states involving CC (II). (II) is useful for generating antibodies against it, detecting or CC quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical CC diagnost of sites expressing (II). (I) and (II) are useful for treating CC disorders involving aberrant protein expression or biological activity. CC The polypeptide and polynucleotide sequences have applications in CC responsible for genetic disorders or other traits to assess biodiversity and the responsible for genetic disorders or other traits to assess biodiversity and contact demonstrate demonstrate and the process of farts and resolute traits to assess biodiversity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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at ftp.wipo.int/pub/published_pct_
                                                              and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic amino acid sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAR-2000; 2000US-0540217
23-AUG-2000; 2000US-0649167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-MAR-2001; 2001WO-US08631.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human diagnostic protein #22869.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABG22878 standard; Protein; 189.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chromosome mapping; gene mapping; gene therapy; forensic
upplement; medical imaging; diagnostic; genetic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 MPLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
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                 sequence data for this patent did not appear in the tion, but was obtained in electronic format directly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID No 53237; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ilarity 100.0%;
Conservative (
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; Mismatches 0;
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from WIPC
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Qγ

21

LPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQASAGPLIGEEAREVARRQEGA

Matches Query Match Best Local :

Similarity

83.2%; ilarity 100.0%; Conservative

0;

Score 461; DB 19; Pred. No. 1.3e-43; 0; Mismatches 0;

Length Indels

0,

Gaps

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RESULT 8
AAW44044
ID AAW4
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Best Local S
Matches 103
                               This sequence represents the peptide hCS-85. It is a peptide of the invention, and has corticostatin or somatostatin activity. Antibodies recognising hCS-17 can be used to screen for a compound that modulates, i.e. an agonist or antagonist, the binding of hCS-17 to its receptor, and to assay for hCS-17, e.g. diagnosis. hCS-17, the DNA encoding it or a receptor agonist or antagonist can be used to treat and prevent hormone-producing tumours (e.g. tumours producing gastrin or insulin), gastric ulcers and dementia, regulate sleep and control growth disorders (e.g. acromegaly, gigantism and dwarfism). They can also be used to regulate hormone secretion and the digestive system (e.g. to treat
                                                                                                                                                                                                                                                                                                                           15-OCT-1996;
07-JUN-1996;
19-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Corticostatin; somatostatin; hCS-85 peptide; hormone-producing gastric ulcer; dementia; growth disorder; hormone secretion red digestive system regulation; neural inhibitor; therapy.
Sequence
                     diabetes),
                                                                                                                                                                 Claim 7; Page 115; 174pp;
                                                                                                                                                                                          growth abnormalities
                                                                                                                                                                                                      anticancer and
                                                                                                                                                                                                                                        N-PSDB; AAV02052, AAV02053.
                                                                                                                                                                                                                                                      WPI; 1998-042177/04.
                                                                                                                                                                                                                                                                             Fukusumi S,
                                                                                                                                                                                                                                                                                                                                                                           05-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
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                                                                                                                                                                                                                Peptide having
                                                                                                                                                                                                                                                                                                   (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hCS-85 peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-JUN-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
85
                      and as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                             Hinuma
A,
                                                                                                                                                                                                                                                                                                                           96JP-0272422.
96JP-0146052.
96JP-0247710.
                                                                                                                                                                                                               corticostatin
                                                                                                                                                                                                      antiulcer
                                                                                                                                                                                                                                                                                                                                                                           97WO-JP01911
                     tumour multiplication or neural inhibitors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peptide;
                                                                                                                                                                                                                                                                            ŝ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96.7%;
97.2%;
                                                                                                                                                                                                                                                                             Kitada
                                                                                                                                                                                                    atin or agent,
                                                                                                                                                                  Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 535.5; DB 2
Pred. No. 1.8e-51;
                                                                                                                                                                                                                                                                            Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A
                                                                                                                                                                                                     somatostatin activity - useful as and for control of dementia and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                        and
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AAM44056
ID AAM4
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AC AAM4
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DT 26-J
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KW Gast
KW Gast
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                                                                                                         Matches
                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                  This sequence is a mutant hCS peptide. It is a peptide of the invention, and has corticostatin or somatostatin activity. Antibodies recognising hCS-17 can be used to screen for a compound that modulates, i.e. an agonist or antagonist, the binding of hCS-17 to its receptor, and to assay for hCS-17, e.g. diagnosis. hCS-17, the DNA encoding it or a receptor agonist or antagonist can be used to treat and prevent hormone-producing tumours (e.g. tumours producing gastrin or insulin), gastric ulcers and dementia, regulate sleep and control growth disorders (e.g. acromegaly, gigantism and dwarfism). They can also be used to regulate hormone secretion and the digestive system (e.g. to treat diabetes), and as tumour multiplication or neural inhibitors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-OCT-1996;
07-JUN-1996;
19-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Corticostatin; somatostatin; hCS peptide; hormone-produ gastric ulcer; dementia; growth disorder; hormone secre digestive system regulation; neural inhibitor; therapy.
                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 118;
                                                                                                                                                                                                                                                                                                                                                                                        Peptide having anticancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                             growth abnormalities
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fukusumi S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TAKE ) TAKEDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-DEC-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW44056 standard;
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                                                                                                         88;
                                                                                                                                                                                                                                                                                                                                                                                                                               AAV02096,
SAGPLIGEEAREVARRQEGAPPQQSARR
               SAGPLIGEEAREVARRQEGAPPQQSARR
                                                  MPLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PPQQSARRDRMPCRNFFWKTFSSCK 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PPQQSARRDRMPCRNFFWKTFSSCK 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQASAGPLIGEEAREVARRQEGA 60
                                                                                                                                                             88 AA;
                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hinuma S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96JP-0272422.
96JP-0146052.
96JP-0247710.
                                                                                                                                                                                                                                                                                                                                                                                      corticostatin or somatostatin activity - useful as antiulcer agent, and for control of dementia and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CHEM IND LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                               AAV02097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide;
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                                                                                                                    81.2%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                174pp; Japanese.
                                                                                                    Score 450; DB 17, Pred. No. 2.3e-42;
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88
                       88
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                                                                                                                              Length 88
                                                                                                      Indels.
                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tumour;
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RESULT 11
AAW42033
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AAW44043
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                                                                                                                                                                                                                        Query Match
Best Local S
Matches 62
                                                                                                                                                                                                                                                                                                  This sequence represents the peptide hCS-62. It is a peptide of the invention, and has corticostatin or somatostatin activity. Antibodies recognising hCS-17 can be used to screen for a compound that modulates, i.e. an agonist or antagonist, the binding of hCS-17 to its receptor, and to assay for hCS-17, e.g. diagnosis. hCS-17, the DNA encoding it or a receptor agonist or antagonist can be used to treat and prevent hormone-producing tumours (e.g. tumours producing gastrin or insulin), gastric ulcers and dementia, regulate sleep and control growth disorders (e.g. acromegaly, gigantism and dwarfism). They can also be used to regulate hormone secretion and the digestive system (e.g. to treat diabetes), and as tumour multiplication or neural inhibitors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-OCT-1996;
07-JUN-1996;
19-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Corticostatin; somatostatin; hCS-62 peptide; hormone-producing tumour; gastric ulcer; dementia; growth disorder; hormone secretion regulation digestive system regulation; neural inhibitor; therapy.
                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                   AAW42033 standard; Protein; 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       growth abnormalities
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide having corticostatin 
anticancer and antiulcer ager
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fukusumi S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-DEC-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW44043;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (TAKE ) TAKEDA CHEM IND LTD.
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DB; AAV02050, AF
                                                                                                                                            104
                                                                                                                                                                                                                                                                                                                                                                                                                                                              7;
                                                                                                                                                                                   44 SSLLTFLAWWFEWTSQASAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSS
                                                                                                                   61
                                                                                                                                                                      \vdash
                                                                                                                                                                                                                        Similarity
62; Conserv
                                                                                                                   CK 62
                                                                                                                                          CK 105
                                                                                                                                                                       SSLLTFLAWWFEWTSQASAGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page
                                                                                                                                                                                                                                                                             62
                                                                                                                                                                                                                       61.4%;
ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hinuma S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                             ΑĄ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             114; 174pp;
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96JP-0146052.
96JP-0247710.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kitada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  atin or somatostatin activity - useful as agent, and for control of dementia and
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          62
                                                                                                                                                                                                                         Score 340; DB 19;
Pred. No. 2.5e-30;
0; Mismatches 0;
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                                                                                                                                                                                                                                                 Length 62
                                                                                                                                                                                                                         Indels
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AAW42033;

28-MAY-1998

(first entry)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This amino acid sequence is the novel rat preprocortistatin protein. CC Cleavage at the N-terminal signal peptide site leads to the formation of procortistatin, which is processed at any of the tandem basic amino cC acid pairs Lys-Lys, or Lys-Arg to produce the mature cortistatin of proteins referred as cortistatin-29, and cortistatin-14. The purified cortistatin, and its agonists, are used to induce sleep while its receptor antagonists (particularly antibodies) is used to inhibit sleep. Although cortistatin is structurally similar to somatostatin, it is able ct depress neuronal electrical activity, induce low frequency waves in the cerebral cortex, antagonise acetylcholine and therefor enhance slow-wave sleep. The antibodies, and oligonucleotide primers, are used in usual immunoassays and hybridisation/amplification assays to detect cor quantify cortistatin (including that administered therapeutically) or its nucleic acid. Oligonucleotides, e.g. antisense molecules, are used in the cortistatin gene may provide diagnosis of sleep-related or nutration related disorders or diseases of the brain.
                                                                                                                      Query Match
Best Local S
Matches 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rat preprocortistatin; N-terminal signal peptide; procortistatin; cortistatin-14; agonist; antibody; inhibition; sleep; somatostatin; neuronal electrical activity; cerebral cortex; antagonist; acetylcholine; primer; PCR; immunoassay; hybridisation; amplification.
                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 102; 128pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New cortistatin peptide(s) - used to mutation(s) and screen for drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SCRI ) SCRIPPS RES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rattus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1998-008886/01.
 69
                               62
                                                                                          N
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                                                                                                                                   Similarity
             AGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                      PSALSLLLLLLSGIAASALPLESGPTGQDS---VQDATGGRRTGLLTFLAWWHEWASQDS
                                                                          PLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQAS
SSTAFEGGTPELSKRQERPPLQQPPHRDKKPCKNFFWKTFSSCK 112
                                                                                                                                                                               112
                                                                                                                      Conservative
                                                                                                                                                                               AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96US-0648322
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81..82
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1..27
                                                                                                                                  53.2%; Score 295; 55.8%; Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "cleaves to give rise residue peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "cleaves to give a putative 29-residue precursor"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "mature peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Siggins
                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modulate sleep, detect
                                                                                                                  DB 19;
4.9e-25;
hes 31;
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                                                                                                                                              Length 112;
                                                                                                                    Indels
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                                                                                                                    2
                                                                                                                    Gaps
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AAEZULT 12
AAEZULT 12
AAEZULT 0
ID AAEZULT 12
XX AAEZULT 16-JU
XX Rat;
KW Rat;
KW Sleep
XX Rattu
XX PT Pepti
FT Prote
FT Cleav
FT Regio
FT Cleav
FT Regio
XX US200
XX IS-MA
XX
                                                                                                  Query Match
Best Local S
Matches 58
                                                                                                                                                                                                                                                                      disorders. Cortistatin proteins may be used as immunogen to produce antibodies immunoreactive with cortistatin, in in vitro ligand binding specificities, to characterise candidate pharmaceutical compounds useful for modulating cortistatin function, and as therapeutic agents for effecting cortistatin functions. The present sequence is rat
                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a substantially isolated and purified mammalian cortistatin. The cortistatin nucleic acids, proteins, polypeptides and antibodies are useful in screening and diagnostic methods, and in therapeutic methods related to modulation of sleep and sleeping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; Fig 1; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New mammalian cortistatin useful in screening and diagnostic methods, and in the rapeutic methods related in modulating sleep and sleeping disorders \,\cdot\,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-MAY-1997;
15-MAY-1996;
                                                                                                                                                                                                                                                 preprocortistatin protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SCRI ) SCRIPPS RES INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2002-328475/36.
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p disorder; pl
11
                                                 N
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                                                                                                                          Similarity
PSALSLLLLLLSGIAASALPLESGPTGQDS---VQDATGGRRTGLLTFLAWWHEWASQDS
                                                 PLSPGLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQAS
                                                                                                                                                                                                     112
                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001US-0766396
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                                                                                                                                                                                                        A
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96US-0648322
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97..98
99..112
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82..83
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sedative;
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                                                                                                                          53.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Peptide formed by proteolytic cleavage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Peptide formed by proteolytic cleavage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mature_cortistatin_protein
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                                                                                                  13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neuropeptide; screening; therapeutic;
cal compound; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       112
                                                                                                Score 295; DB 23;
Pred. No. 4.9e-25;
3; Mismatches 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A
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                                                                                                  31;
                                                                                                                                                 Length
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62

AGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105

밁

69

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RESULT 1:
AAW42034
ID AAWW
XX
AC AAW
XX
AC AAW
XX
DT 28-1
DT 28-1
DT W
COI
KW COI
KW Som
This amino acid sequence is the novel mouse preprocortistatin protein. CC Cleavage at the N-terminal signal peptide site leads to the formation CC of procortistatin, which is processed at any of the tandem basic amino CC acid pairs Lys-Ser, or Lys-Lys to produce the mature cortistatin CC proteins referred as mouse cortistatin-29, and cortistatin-14. The purified cortistatin, and its agonists, are used to induce sleep while CC its receptor antagonists (particularly antibodies) is used to inhibit CC sleep. Although cortistatin is structurally similar to somatostatin, CC sleep. Although cortistatin is structurally similar to somatostatin, CC it is able to depress neuronal electrical activity, induce low frequency waves in the cerebral cortex, antagonise acetylcholine and therefor CC enhance slow-wave sleep. The antibodies, and oligonucleotide primers, CC are used in usual immunoassays and hybridisation/amplification assays to CC detect or quantify cortistatin (including that administered complecules, are used in vivo to acid. Oligonucleotides, e.g. antisense CC molecules, are used in vivo to cortistatin gene may provide diagnosis of sleep-related or neuronal depression-related disorders or diseases of
                               sleep-related
the brain.
                                                                                                                                                                                                                                                                                                                                                            Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                         New cortistatin peptide(s) - used mutation(s) and screen for drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              De Lecea L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cleavage-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                            1998-008886/01
DB; AAV09156.
                                                                                                                                                                                                                                                                                                                                                            Page 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Henriksen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
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/note=
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79..80
/note-
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1..27
                                                                                                                                                                                                                                                                                                                                                        128pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "cleaves to give a precursor"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "cleaves to give rise 
14 residue peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "mature peptide"
                                                                                                                                                                                                                                                                                                                                                        English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Siggins GR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         109
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                                                                                                                                                                                                                                                                                                                                                                                                           sleep,
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Sequence

109

A

preprocortistatin

protein

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RESULT 14
AAE21872
ID AAE2:
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Best Local S
Matches 58
       The invention relates to a substantially isolated and purified mammalian cortistatin. The cortistatin nucleic acids, proteins, polypeptides and antibodies are useful in screening and diagnostic methods, and in therapeutic methods related to modulation of sleep and sleeping disorders. Cortistatin proteins may be used as immunogen to produce antibodies immunoreactive with cortistatin, in in vitro ligand binding specificities, to characterise candidate pharmaceutical compounds useful for modulating cortistatin functions. The present sequence is mouse effecting cortistatin functions. The present sequence is
                                                                                                                                                                                                       WPI; 200
N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cleavage-
Region
                                                                                                                               Claim 2; Fig 3; 50pp; English.
                                                                                                                                                                  and in therapeutic
                                                                                                                                                                                                                                           Sutcliffe JG,
                                                                                                                                                                                                                                                                                          15-MAY-1997;
15-MAY-1996;
                                                                                                                                                                                                                                                                                                                              18-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                      31-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                               Cleavage-site
Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sleep disorder;
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                                                                                                                                                                     New mammalian cortistatin useful in and in therapeutic methods related i
                                                                                                                                                                                                                                                                   (SCRI ) SCRIPPS RES INST
                                                                                                                                                                                                                                                                                                                                                                            US2002013456-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse preprocortistatin protein.
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                                                                                                                                                                                                         2002-328475/36
DB; AAD34525.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 PLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQAS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SSTPVGGGTPGLSKSQERPPPQQPPHLDKKPCKNFFWKTFSSCK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                              2001US-0766396
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                                                                                                                                                                                                                                           Lecea
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96US-0648322
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26..109
/label=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        :in; sedative; neuropeptide; screening; therapeutic;
pharmaceutical compound; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers 1..25
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                                                                                                                                                                                                                                                                                                                                                                                                                109
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                                                                                                                                                                                                                                                                                                                                                                                                    "Peptide formed
                                                                                                                                                                                                                                                                                                                                                                                                                                       "Peptide formed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mature_cortistatin_protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Signal_peptide
                                                                                                                                                                                                                                            Henriksen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 272; DB 19;
Pred. No. 1.7e-22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B
                                                                                                                                                                   in screening and diagnostic methods,
in modulating sleep and sleeping
                                                                                                                                                                                                                                           SJ,
                                                                                                                                                                                                                                                                                                                                                                                                   bу
                                                                                                                                                                                                                                                                                                                                                                                                                                       by proteolytic cleavage"
                                                                                                                                                                                                                                           Siggins GR;
                                                                                                                                                                                                                                                                                                                                                                                                    proteolytic cleavage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 109
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RESULT 15
AAW42035
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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                                                                               This amino acid sequence is the novel rat procortistatin protein, which is cleaved to produce three peptides (AAM42036, AAM42037, AAM42038). One of the peptides (AAM42038), designated as rat cortistatin-14, is highly conserved among species e.g. mouse and human. The purified cortistatin, and its agonists, are used to induce sleep while its receptor antagonists (particularly antibodies) is used to inhibit sleep. Although cortistatin is structurally similar to somatostatin, it is able to depress neuronal electrical activity, induce low frequency waves in the cerebral cortex, antagonise acetylcholine and therefor enhance to depress neuronal electrical activity, induce low frequency waves in the cerebral cortex, antagonise acetylcholine and therefor enhance to slow-wave sleep. The antibodies, and oligonucleotide primers, are used in usual immunoassays and hybridisation/amplification assays to detect or quantify cortistatin (including that administered therapeutically) or its nucleic acid. Oligonucleotides, e.g. antisense molecules, are used in the cortistatin gene may provide diagnosis of sleep-related or neuronal of the cortistatin gene may provide diagnosis of sleep-related or neuronal or the cortistatin gene may provide diagnosis of the brain.
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rat procortistatin; preprocortistatin; cortistatin-29; cortistatin-14; agonist; antibody; inhibition; sleep; somatostatin; acetylcholine; neuronal electrical activity; cerebral cortex; antagonist; primer; PCR; immunoassay; hybridisation; amplification.
                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 105; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      De Lecea L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-MAY-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SCRI ) SCRIPPS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rattus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rat procortistatin protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-MAY-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62 AGPLIGEEAREVARRQEGAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              66 SSTPVGGGTPGLSKSQERPPPQQPPHLDKKPCKNFFWKTFSSCK 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 PLSPGLLLLLLSGATATAALPLEGGPTGRDSEHMQEAAGIRKSSLLTFLAWWFEWTSQAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PSAFGLLLL---WGVAASALPLESGPTGQDS--VQEATEGR-SGLLTFLAWWHEWASQAS
                                                     85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            109 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Henriksen SJ,
                                                   AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96US-0648322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97WO-US08481.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Protein;
48.2%;
57.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    49.18; 55.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Siggins GR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 272; DB 23;
Pred. No. 1.7e-22;
11; Mismatches 29;
Score 267; DB 19;
Pred. No. 4.5e-22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sutcliffe
            Length 85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61
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Qy
  밁
                                     В
                                                       Q
                                                                            Matches
59 RPPLQQPPHRDKKPCKNFFWKTFSSCK 85
           79 GAPPQQSARRDRMPCRNFFWKTFSSCK 105
                                               50;
                                      SALPLESGPTGQDS---VQDATGGRRTGLLTFLAWWHEWASQDSSSTAFEGGTPELSKRQE
                                                                            Conservative
                                                                            12;
                                                                            Mismatches
                                                                            23;
                                                                            Indels
                                                                            ?
                                                                           Gaps
                                      58
                                                         78
```

Search completed: June 23, 2003, 15:54:43 Job time: 76 secs

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